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Figure 1: construction of a synthetic human antibody library based on consensus sequences

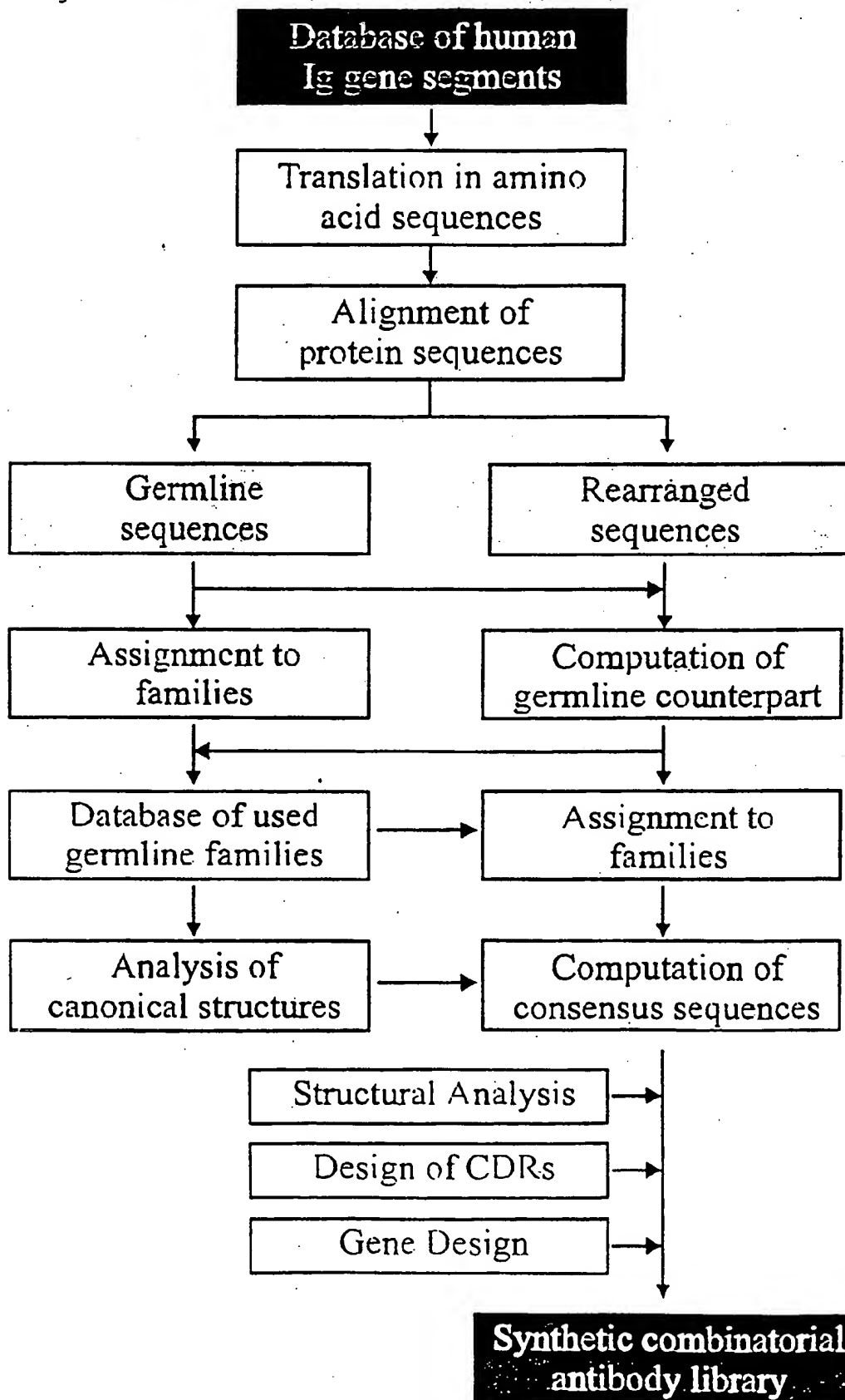


Figure 2A: VL kappa consensus sequences

framework 1														CDRI																
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30
Vκ1	D	I	Q	M	T	Q	S	P	S	S	L	S	A	S	V	G	D	R	V	T	I	T	C	R	A	S	Q	-	-	-
Vκ2	D	I	V	M	T	Q	S	P	L	S	L	P	V	T	P	G	E	P	A	S	I	S	C	R	S	S	Q	S	L	L
Vκ3	D	I	V	L	T	Q	S	P	A	T	L	S	L	S	P	G	E	R	A	T	L	S	C	R	A	S	Q	S	-	-
Vκ4	D	I	V	M	T	Q	S	P	D	S	L	A	V	S	L	G	E	R	A	T	I	N	C	R	S	S	Q	S	V	L

CDRI										framework 2														CDR II						
D	F	F	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	
Vκ1	-	-	-	G	I	S	S	Y	L	A	W	Y	Q	Q	K	P	G	K	A	P	K	L	L	I	Y	A	A	S	S	L
Vκ2	H	S	-	N	G	Y	N	Y	L	D	W	Y	L	Q	K	P	G	Q	S	P	Q	L	L	I	Y	L	G	S	N	R
Vκ3	-	-	-	V	S	S	Y	L	A	W	Y	Q	Q	K	P	G	Q	A	P	R	L	L	I	Y	G	A	S	S	R	
Vκ4	Y	S	S	N	N	K	N	Y	L	A	W	Y	Q	Q	K	P	G	Q	P	P	K	L	L	I	Y	W	A	S	T	R

[illegible][illegible]

Figure 2B: VL lambda consensus sequences

framework 1														CDRI																
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28			
VA1	Q	S	V	L	T	Q	P	P	S	-	V	S	G	A	P	G	Q	R	V	T	I	S	C	S	G	S	S	N	I	
VA2	Q	S	A	L	T	Q	P	A	S	-	V	S	G	S	P	G	Q	S	I	T	I	S	C	T	G	T	S	S	D	V
VA3	S	Y	E	L	T	Q	P	P	S	-	V	S	V	A	P	G	Q	T	A	R	I	S	C	S	G	D	A	-	-	L

CDRI				framework 2														CDRI II													
29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57			
VA1	G	S	N	-	Y	V	S	W	Y	Q	Q	L	P	G	T	A	P	K	L	I	Y	D	N	N	Q	R	P	S	G		
VA2	G	G	Y	N	Y	V	S	W	Y	Q	Q	H	P	G	K	A	P	K	L	M	I	Y	D	V	S	N	R	P	S	G	
VA3	G	D	K	-	Y	A	S	W	Y	Q	Q	K	P	G	Q	A	P	V	L	V	I	Y	D	D	S	D	R	P	S	G	

Figure 2B: VL lambda consensus sequences

framework 3	
VA1	V P D R F S G S K S G T S A S L A I T G L Q S E D E A D Y Y
VA2	V S N R F S G S K S G N T A S L T I S G L Q A E D E A D Y Y
VA3	I P E R F S G S N S G N T A T L T I S G T Q A E D E A D Y Y
framework 4	
CDRIII	
VA1	C Q Q H Y T T P P V F G G G T K L T V L G
VA2	C Q Q H Y T T P P V F G G G T K L T V L G
VA3	C Q Q H Y T T P P V F G G G T K L T V L G

Figure 2C: V heavy chain consensus sequences

framework 1																														
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30
VH1A	Q	V	Q	L	V	Q	S	G	A	E	V	K	K	P	G	S	S	V	K	V	S	C	K	A	S	G	G	T	F	S
VH1B	Q	V	Q	L	V	Q	S	G	A	E	V	K	K	P	G	A	S	V	K	V	S	C	K	A	S	G	G	T	F	T
VH2	Q	V	Q	L	K	E	S	G	P	A	L	V	K	P	T	Q	T	L	T	L	T	C	T	F	S	G	F	S	L	S
VH3	E	V	Q	L	V	E	S	G	G	L	V	Q	P	G	S	L	R	L	S	C	A	A	S	G	F	T	F	S		
VH4	Q	V	Q	L	Q	E	S	G	P	G	L	V	K	P	S	E	T	L	S	L	T	C	T	V	S	G	G	S	I	S
VH5	E	V	Q	L	V	Q	S	G	A	E	V	K	K	P	G	E	S	L	K	I	S	C	K	G	S	G	Y	S	F	T
VH6	Q	V	Q	L	Q	Q	S	G	P	G	L	V	K	P	S	Q	T	L	S	L	T	C	A	I	S	G	D	S	V	S

CDRI		framework 2																CDR II																
	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57							
VH1A	S	-	-	Y	A	I	S	W	V	R	Q	A	P	G	Q	G	L	E	W	M	G	G	I	I	P	-	-	-	I	F	G	T	A	
VH1B	S	-	-	Y	Y	M	H	W	V	R	Q	A	P	G	Q	G	L	E	W	M	G	W	I	N	P	-	-	-	-	N	S	G	G	T
VH2	T	S	G	V	G	V	G	W	I	R	Q	P	P	G	K	A	L	E	W	L	A	L	I	D	-	-	-	-	-	W	D	D	D	K
VH3	S	-	-	Y	A	M	S	W	V	R	Q	A	P	G	K	G	L	E	W	V	S	A	I	S	G	-	-	-	-	S	G	G	S	T
VH4	S	-	-	Y	Y	W	S	W	I	R	Q	P	P	G	K	G	L	E	W	I	G	Y	I	Y	-	-	-	-	-	Y	S	G	S	T
VH5	S	-	-	Y	W	I	G	W	V	R	Q	M	P	G	K	G	L	E	W	M	G	I	I	Y	P	-	-	-	-	G	D	S	D	T
VH6	S	N	S	A	A	W	N	W	I	R	Q	S	P	G	R	G	L	E	W	L	G	R	T	Y	Y	R	-	-	-	S	K	W	Y	N

Figure 2C: V heavy chain consensus sequences.

CDRII		framework 3	
VH1A	58	N	85
VH1B	59	N	84
VH2	60	Y	83
VH3	61	Y	82
VH4	62	N	81
VH5	63	R	80
VH6	64	D	79
	65	Y	78
	66	S	77
	67	T	76
	68	S	75
	69	E	74
	70	A	73
	71	D	72
	72	T	71
	73	R	70
	74	D	69
	75	T	68
	76	S	67
	77	I	66
	78	T	65
	79	M	64
	80	T	63
	81	I	62
	82	S	61
	83	V	60
	84	K	59
	85	G	58
	86	R	57
	87	V	56
	88	F	55
	89	Q	54
	90	K	53
	91	S	52
	92	V	51
	93	L	50
	94	K	49
	95	S	48
	96	V	47
	97	P	46
	98	S	45
	99	F	44
	100	Q	43
	101	D	42
	102	T	41
	103	S	40
	104	K	39
	105	N	38
	106	P	37
	107	S	36
	108	V	35
	109	L	34
	110	Q	33
	111	F	32
	112	T	31
	113	S	30
	114	V	29
	115	T	28
	116	V	27
	117	T	26
	118	V	25
	119	T	24
	120	V	23
	121	T	22
	122	V	21
	123	T	20
	124	V	19
	125	T	18
	126	V	17
	127	T	16
	128	V	15
	129	T	14
	130	V	13
	131	T	12
	132	V	11
	133	T	10
	134	V	9
	135	T	8
	136	V	7
	137	T	6
	138	V	5
	139	T	4
	140	V	3
	141	T	2
	142	V	1

Figure 3A: V kappa 1 (Vk1) gene sequence

```

.D I Q M T Q S P S S L S A S V G D
EcoRV
~~~~~
BanII
~~~~~
GATATCCAGA TGACCCAGAG CCCGTCTAGC CTGAGCGCGA GCGTGGGTGA
CTATAGGTCT ACTGGGTCTC GGCAGATCG GACTCGCGCT CGCACCCACT

R V T I T C R A S Q G I S S Y L
PstI
~~~~~
TCGTGTGACC ATTACCTGCA GAGCGAGCCA GGCATTAGC AGCTATCTGG
AGCACACTGG TAATGGACGT CTCGCTCGGT CCCGTAATCG TCGATAGACC

A W Y Q Q K P G K A P K L L I Y A
KpnI SexAI AseI
~~~~~
CGTGGTACCA GCAGAAACCA GGTAAGCAC CGAAACTATT AATTATGCA
GCACCATGGT CGTCTTTGGT CCATTTCGTG GCTTTGATAA TTAATAACGT

A S S L Q S G V P S R F S G S
SanDI BamHI
~~~~~
GCCAGCAGCT TGCAAAGCGG GGTCCTCGTCC CGTTTATAGC GCTCTGGATC

```

Figure 3A: V kappa 1 (Vκ1) gene sequence (continued)

CGGTCGTCGA ACGTTTCGCC CCAGGGCAGG GCAAATCGC CGAGACCTAG

G T D F T L T I S S L Q P E D F

Eco57I

~~~~~

BamHI

BbsI

~~~~~

CGGCACTGAT TTTACCCTGA CCATTAGCAG CCTGCAACCT GAAGACTTTG

GCCGTGACTA AAATGGGACT GGTAATCGTC GGACGTTGGA CTTCTGAAAC

A T Y Y C Q Q H Y T T P P T F G Q

MscI

~~~~~

CGACCTATTA TTGCCAGCAG CATTATACCA CCCC GCCGAC CTTTGGCCAG

GCTGGATAAT AACGGTCGTC GTAATATGGT GGGCGGCTG GAAACCGGTC

G T K V E I K R T

BsiWI

~~~~~

GGTACGAAAG TTGAAATTAA ACGTACG

CCATGCTTTC AACTTAATT TGCATGC

Figure 3B: V kappa 2 (Vk2) gene sequence

```

D I V M T Q S P L S L P V T P G E
EcoRV          BanII
~~~~~
GATATCGTGA TGACCCAGAG CCCACTGAGC CTGCCAGTGA CTCCGGGGCGA
CTATAGCACT ACTGGGTCTC GGGTGACTCG GACGGTCACT GAGCCCCCGCT

P A S I S C R S S Q S L L H S N
PstI
~~~~~
GCCTGCGAGC ATTAGCTGCA GAAGCAGCCA AAGCCTGCTG CATAGCAACG
CGGACGCTCG TAATCGACGT CTTCGTCGGT TTCGGACGAC GTATCGTTGC

G Y N Y L D W Y L Q K P G Q S P Q
KpnI          SexAI
~~~~~
GCTATAACTA TCTGGATTGG TACCTTCAA AACCAGGTCA AAGCCCCGAG
CGATATTGAT AGACCTAACC ATGGAAGTTT TTGGTCCAGT TTCGGGCGTC

L L I Y L G S N R A S G V P D R F
AseI          SmaI
~~~~~
CTATTAATT ATCTGGGCAG CAACCGTGCC AGTGGGGTCC CGGATCGTTT
GATAATTAAA TAGACCCGTC GTTGGCACGG TCACCCAGG GCCTAGCAAA

```

Figure 3B: V kappa 2 (Vk2) gene sequence (continued)

S	G	S	G	S	G	T	D	F	T	L	K	I	S	R	V
BamHI															
~~~~~															
T	A	G	C	G	G	C	T	C	G	G	C	A	A	A	T
A	T	C	G	C	C	G	A	G	A	A	T	G	G	A	C
~~~~~															
E	A	E	D	V	G	V	Y	Y	C	Q	Q	H	Y	T	P
Eco57I															
~~~~~															
BbsI															
~~~~~															
A	A	G	C	T	G	A	A	G	A	G	A	A	T	T	A
T	T	C	G	A	C	T	T	C	T	G	C	G	T	A	A
~~~~~															
P	T	F	G	Q	G	T	K	V	E	I	K	R	T		
MscI															
~~~~~															
C	C	G	A	C	C	T	T	G	G	C	C	G	T	A	A
G	G	T	G	G	A	A	C	A	A	T	T	G	C	A	T
~~~~~															
BsiWI															
~~~~~															
C	C	G	A	C	C	T	T	G	G	A	A	T	T	G	C
G	G	T	G	G	A	A	C	A	A	T	T	G	C	A	T

Figure 3C: V kappa 3 (V κ 3) gene sequence

```

D I V L T Q S P A T L S L S P G E
EcoRV                               BanII
~~~~~
GATATCGTGC TGACCCAGAG CCCGGCGACC CTGAGCCTGT CTCCGGGCGA
CTATAGCAGG ACTGGGTCTC GGGCCGCTGG GACTCGGACA GAGGCCCGCT

R A T L S C R A S Q S V S S Y
PstI
~~~~~
ACGTGCGACC CTGAGCTGCA GAGCGAGCCA GAGCGTGAGC AGCAGCTATC
TGCACGCTGG GACTCGACGT CTCGCTCGGT CTCGCACTCG TCGTCGATAG

L A W Y Q Q K P G Q A P R L L I Y
KpnI                               SexAI                               AseI
~~~~~
TGGCGTGGTA CCAGCAGAAA CCAGGTCAAG CACCGCGTCT ATTAATTAT
ACCGCACCAT GGTCTCTTT GTCCAGTTC GTGGCGCAGA TAATTAAATA

G A S S R A T G V P A R F S G S G
                               SandI                               BamHI
~~~~~
GGCGCGAGCA GCCGTGCAAC TGGGGTCCCG GCGCGTTTTA GCGGCTCTGG

```

Figure 3C: V kappa 3 (V κ 3) gene sequence (continued)

CCGCGGCTCGT CGGCACGTTG ACCCCAGGGC CGCGCAAAAT CGCCGAGACC

S G T D F T L T I S S L E P E D
Eco57I
~~~~~

BamHI  
~~~~~  
ATCCGGCAGG GATTTACCC TGACCATTAG CAGCCTGGAA CCTGAAGACT
TAGGCCGTGC CTAAAATGGG ACTGGTAATC GTCGGACCTT GGA CTCTGA

F A V Y Y C Q Q H Y T T P P T F G
MscI
~~~~~

TTGCGGTGTA TTATTGCCAG CAGCATTATA CCACCCCGCC GACCTTTGGC  
AACGCCACAT AATAACGGTC GTCGTAATAT GTGGGGCGG CTGGAACCCG

Q G T K V E I K R T  
MscI  
~~~~~  
CAGGTACGA AAGTTGAAAT TAAACGTACG
GTCCCATGCT TTCAACTTA ATTGCATGC

Figure 3D: V kappa 4 (Vκ4) gene sequence

```

D I V M T Q S P D S L A V S L G E
EcoRV      BanII
~~~~~
GATATCGTGA TGACCCAGAG CCCGGATAGC CTGGCGGTGA GCCTGGGCGGA
CTATAGCACT ACTGGGTCTC GGGCCTATCG GACCGCCACT CGGACCCGCT

R A T I N C R S S Q S V L Y S S
PstI
~~~~~
ACGTGCGACC ATTAAGTGA GAAGCAGCCA GAGCGTGCTG TATAGCAGCA
TGCACGCTGG TAATTGACGT CTTCGTCGGT CTCGCACGAC ATATCGTCGT

N N K N Y L A W Y Q Q K P G Q P P
KpnI      SexAI
~~~~~
ACAACAAAAA CTATCTGGCG TGTACCAGC AGAAACCAGG TCAGCCGCCG
TGTGTTTGT GATAGACCGC ACCATGGTCG TCTTTGGTCC AGTCGGCGGC

K L L I Y W A S T R E S G V P D R
AseI      SandI
~~~~~
AAACTATTAA TTTATTGGGC ATCCACCCGT GAAAGCGGGG TCCCGGATCG
TTTGATAATT AAATAACCCG TAGGTGGGCA CTTTCGCCCC AGGCCTAGC

```

Figure 3D: V kappa 4 (Vκ4) gene sequence (continued)

```

      F  S  G  S  G  S  G  T  D  F  T  L  T  I  S  S
          BamHI
      ~~~~~
      TTTTAGCGGC TCTGGATCCG GCACTGATTT TACCCCTGACC ATTTCGTCCC
      AAAATCGCCG AGACCTAGGC CGTGACTAAA ATGGGACTGG TAAAGCAGGG

      L  Q  A  E  D  V  A  V  Y  Y  C  Q  Q  H  Y  T  T
          Eco57I
      ~~~~~
          BbsI
      ~~~~~
      TGCAAGCTGA AGACGTGGCG GTGTATTATT GCCAGCAGCA TTATACCACC
      ACGTTCGACT TCTGCACCGC CACATAATAA CGGTCGTCGT AATATGGTGG

      P  P  T  F  G  Q  G  T  K  V  E  I  K  R  T
          MscI
      ~~~~~
      CCGCCGACCT TTGGCCAGGG TACGAAAGTT GAAATTAAAC GTACG
      GGCGGCTGGA AACCGGTCCC ATGCTTTCAA CTTTAATTG CATGC
  
```

Figure 4A: V lambda 1 (Vλ1) gene sequence

Q S V L T Q P P S V S G A P G Q R
SexAI

CAGAGCGTGC TGACCCAGCC GCCTTCAGTG AGTGGCGCAC CAGTTCAGCG
GTCTCGCACG ACTGGGTCGG CGGAAGTCAC TCACCCGCGTG GTCCAGTCGC
Eco57I

V T I S C S G S S S N I G S N Y
BssSI

TGTGACCATC TCGTGAGCG GCAGCAGCAG CAACATTGGC AGCAACTATG
ACACTGGTAG AGCACATCGC CGTCGTCGTC GTTGTAAACCG TCGTTGATAC

V S W Y Q Q L P G T A P K L L I Y
KpnI XmaI BbeI

TGAGCTGGTA CCAGCAGTTG CCCGGGACGG CGCCGAAACT GCTGATTAT
ACTCGACCAT GGTGTC AAC GGGCCCTGCC GCGGCTTGA CGACTAAATA

D N N Q R P S G V P D R F S G S K
Bsu36I BamHI

Figure 4A: V lambda 1 (Vλ1) gene sequence (continued)

GATAACAACC AGCGTCCCTC AGGCGTGCCG GATCGTTTA GCGGATCCAA
CTATTGTTGG TCGCAGGGAG TCCGCACGGC CTAGCAAAAT CGCCTAGGTT

S G T S A S L A I T G L Q S E D
BbsI

AAGCGGACC AGCGGAGCC TTGCGATTAC GGGCCTGCAA AGCGAAGACG
TTCGCCGTGG TCGCGCTCGG AACGCTAATG CCCGGACGTT TCGCTTCTGC

E A D Y Y C Q Q H Y T T P P V F G
AAGCGGATTA TTATTGCCAG CAGCATTATA CCACCCCGCC TGTGTTTGGC
TTCGCCCTAAT AATAACGGTC GTCGTAATAT GGTGGGGCGG ACACAAACCG

G G T K L T V L G
HpaI MscI
~~~~~  
GGCGGCACGA AGTTAACCGT TCTTGGC  
CCGCCGTGCT TCAATTGGCA AGAACCG

Figure 4B: V lambda 2 (Vλ2) gene sequence

|            |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
|------------|------------|------------|------------|------------|---|---|---|---|---|---|---|---|---|---|---|---|
| Q          | S          | A          | L          | T          | Q | P | A | S | V | S | G | S | P | G | Q | S |
| SexAI      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| CAGAGCGCAC | TGACCCAGCC | AGCTTCAGTG | AGCGGCTCAC | CAGGTCAGAG |   |   |   |   |   |   |   |   |   |   |   |   |
| GTCTCGCGTG | ACTGGGTCGG | TCGAAGTCAC | TCGCCGAGTG | GTCCAGTCTC |   |   |   |   |   |   |   |   |   |   |   |   |
| Eco57I     |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| I          | T          | I          | S          | C          | T | G | T | S | S | D | V | G | G | Y | N |   |
| BssSI      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| CATTACCATC | TCGTGTACGG | GTACTAGCAG | CGATGTGGGC | GGCTATAACT |   |   |   |   |   |   |   |   |   |   |   |   |
| GTAATGGTAG | AGCACATGCC | CATGATCGTC | GCTACACCCG | CCGATATTGA |   |   |   |   |   |   |   |   |   |   |   |   |
| Y          | V          | S          | W          | Y          | Q | Q | H | P | G | K | A | P | K | L | M | I |
| KpnI       |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ATGTGAGCTG | GTACCAGCAG | CATCCCGGGA | AGCGGCCGAA | ACTGATGATT |   |   |   |   |   |   |   |   |   |   |   |   |
| TACACTCGAC | CATGTCGTC  | GTAGGGCCCT | TCCGCGGCTT | TGACTACTAA |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| Y          | D          | V          | S          | N          | R | P | S | G | V | S | N | R | F | S | G | S |
| Bsu36I     |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| TATGATGTGA | GCAACCGTCC | CTCAGGCGTG | AGCAACCGTT | TTAGCGGATC |   |   |   |   |   |   |   |   |   |   |   |   |
| ATACTACACT | CGTTGGCAGG | GAGTCCGCAC | TCGTTGGCAA | AATCGCCTAG |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| BamHI      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |

Figure 4B: V lambda 2 (Vλ2) gene sequence (continued)

```

      K  S  G  N  T  A  S  L  T  I  S  G  L  Q  A  E
      BamHI
      ~
      CAAAGCGC AACACCGCGA GCCTGACCAT TAGCGGCCCTG CAAGCGGAAG
      GTTTTCGCCG TTGTGGCGCT CGGACTGGTA ATCGCCGGAC GTTCGCCCTTC

      D  E  A  D  Y  Y  C  Q  Q  H  Y  T  T  P  P  V  F
      BbsI
      ~
      ACGAAGCGG TATTATTGC CAGCAGCATT ATACCACCCC GCCGTGTGTTT
      TGCTTCGCC TATAATAACG GTCGTCGTAA TATGGTGGGG CGGACACAAA

      G  G  G  T  K  L  T  V  L  G
      HpaI      MscI
      ~~~~~
 GGCGGCGGCA CGAAGTTAAC CGTTCTTGGC
 CCGCCGCCGT GCTTCAATTG GCAAGAACCG

```



Figure 4C: V lambda 3 (Vλ3) gene sequence

|            |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
|------------|------------|------------|------------|------------|---|---|---|---|---|---|---|---|---|---|---|---|
| S          | Y          | E          | L          | T          | Q | P | P | S | V | S | V | A | P | G | Q | T |
| SexAI      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| AGCTATGAAC | TGACCCAGCC | GCCTTCAGTG | AGCGTTGCAC | CAGGTCAGAC |   |   |   |   |   |   |   |   |   |   |   |   |
| TCGATACTTG | ACTGGGTCGG | CGGAAGTCAC | TCGCAACGTG | GTCCAGTCTG |   |   |   |   |   |   |   |   |   |   |   |   |
| Eco57I     |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| A          | R          | I          | S          | C          | S | G | D | A | L | G | D | K | Y | A | S |   |
| BssSI      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| CGCGCGTATC | TCGTGTAGCG | CGGATGCGCT | GGCGGATAAA | TACGCGAGCT |   |   |   |   |   |   |   |   |   |   |   |   |
| GCGCGCATAG | AGCACATCGC | CGCTACGCGA | CCCGCTATTT | ATGCGCTCGA |   |   |   |   |   |   |   |   |   |   |   |   |
| W          | Y          | Q          | Q          | K          | P | G | Q | A | P | V | L | V | I | Y | D | D |
| KpnI       |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| XmaI       |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| BbeI       |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |   |
| GGTACCAGCA | GAAACCCGGG | CAGCGGCCAG | TTCTGGTGAT | TTATGATGAT |   |   |   |   |   |   |   |   |   |   |   |   |
| CCATGGTCGT | CTTTGGGCCC | GTCCGCGGTC | AAGACCACTA | AATACTACTA |   |   |   |   |   |   |   |   |   |   |   |   |

Figure 4C: V lambda 3 (Vλ3) gene sequence (continued)

```

S D R P S G I P E R F S G S N S G
 Bsu36I BamHI
TCTGACCGTC CCTCAGGCAT CCCGGAACGC TTAGCGGAT CCAACAGCGG
AGACTGGCAG GGAGTCCGTA GGCCTTGGC AAATCGCCTA GGTGTGCGCC
      ~~~~~
N T A T L T I S G T Q A E D E A
      BbsI
CAACACCGCG ACCCTGACCA TTAGCGGCAC TCAGCGCGAA GACGAAGCGG
GTTGTGGCGC TGGGACTGGT AATCGCCGTG AGTCCGCCTT CTGCTTCGCC
      ~~~~~
D Y Y C Q Q H Y T P P V F G G G
ATTATTATTG CCAGCAGCAT TATACCACCC CGCCTGTGTT TGGCGGCGGC
TAATAATAAC GGTGTCGTA ATATGTGGG GCGGACACAA ACCGCCGCCG
      ~~~~~
T K L T V L G
      HpaI      MscI
      ~~~~~
ACGAAGTTAA CCGTCTTGG C
TGCTTCAATT GGCAAGAACC G

```

Figure 5A: V heavy chain 1A (VH1A) gene sequence

```

Q V Q L V Q S G A E V K K P G S S
MfeI
~~~~~
CAGGTGCAAT TGGTTCAGTC TGGCGCGGAA GTGAAAAAAC CGGGCAGCAG
GTCCACGTTA ACCAAGTCAG ACCGCGCCTT CACTTTTTCG GCCCGTCGTC

V  K  V  S  C  K  A  S  G  G  T  F  S  S  Y  A
BspEI
~~~~~
CGTGAAAGTG AGCTGCAAAG CCTCCGGAGG CACTTTTAGC AGCTATGCCA
GCACTTTCAC TCGACGTTTC GGAGGCCCTCC GTGAAAATCG TCGATACGCT

I S W V R Q A P G Q G L E W M G G
BstXI XhoI
~~~~~
TTAGCTGGGT GCGCCAAGCC CCTGGGCAGG GTCTCGAGTG GATGGCGCGC
AATCGACCCA CGCGGTTCGG GGACCCGTC CAGAGCTCAC CTACCCGCGC

I  I  P  I  F  G  T  A  N  Y  A  Q  K  F  Q  G  R
ATTATTCCGA TTTTGGCAC GCGGAACACTAC GCGCAGAAGT TTCAGGGCCG
TAATAAGGCT AAAAACCGTG CCGCTTGATG CGCGTCTTCA AAGTCCCGGC

V  T  I  T  A  D  E  S  T  S  T  A  Y  M  E  L
BstEII

```

Figure 5A: V heavy chain 1A (VH1A) gene sequence (continued)

```

~~~~~
GGTGACCATT ACCGCGGATG AAAGCACCAG CACCGCGTAT ATGGAACCTGA
CCACTGGTAA TGGCGCCTAC TTTCGTGGTC GTGGCGCATA TACCTTGACT

S S L R S E D T A V Y Y C A R W G
EagI BssHII
~~~~~
GCAGCCTGCG TAGCGAAGAT ACGCCCGTGT ATTATTGCGC GCGTTGGGCG
CGTCGGACGC ATCGCTTCTA TGCCGGCACA TAATAACGCG CGCAACCCCG

G D G F Y A M D Y W G Q G T L V T
StyI
~~~~~
GGCGATGGCT TTTATGCGAT GGATTATTGG GGCCAAGGCA CCCTGGTGAC
CCGCTACCGA AAATACGCTA CCTAATAACC CCGGTTCCGT GGGACCACTG

V S S
BlnI
~~~~~
GGTAGCTCA G
CCAATCGAGT C

```

Figure 5B: V heavy chain 1B (VH1B) gene sequence

```

Q V Q L V Q S G A E V K K P G A S
MfeI
~~~~~
CAGGTGCAAT TGGTTCAGAG CGGCGCGGAA GTGAAAAAAC CGGGCGCGGAG
GTCCACGTTA ACCAAGTCTC GCCGCGCCTT CACTTTTGTG GCCCGCGCTC

V K V S C K A S G Y T F T S Y Y
BspEI
~~~~~
CGTGAAAGTG AGCTGCAAAG CCTCCGGATA TACCTTTACC AGCTATTATA
GCACTTTCAC TCGACGTTTC GGAGGCCTAT ATGGAAATGG TCGATAATAT

M H W V R Q A P G Q G L E W M G W
BstXI
~~~~~
TGCAC TGGT CCGCCAAGCC CCTGGGCAGG GTCTCGAGTG GATGGGCTGG
ACGTGACCCA GCGGTTTCGG GGACCCGTCC CAGAGCTCAC CTACCCGACC

I N P N S G G T N Y A Q K F Q G R
ATTAAACCCGA ATAGCGGCGG CACGAAC TAC GCGCAGAAGT TTCAGGGCCG
TAATTGGGCT TATCGCCGCC GTGCTTGATG CCGCTCTCA AAGTCCCCGC

```

Figure 5B: V heavy chain 1B (VH1B) gene sequence (continued)

```

V T M T R D T S I S T A Y M E L
BstEII

GGTGACCATG ACCCGTGATA CCAGCATTAG CACCGCGTAT ATGGAAGTGA
CCACTGGTAC TGGGCACTAT GGTCGTAATC GTGGCGCATA TACCTTGACT

S S L R S E D T A V Y Y C A R W G
EagI

BssHII

GCAGCCTGCG TAGCGAAGAT ACGGCCGTGT ATTATTGCGC GCGTTGGGGC
CGTCGGACGC ATCGCTTCTA TGCCGGCACA TAATAACGCG CGCAACCCCG

G D G F Y A M D Y W G Q G T L V T
StyI

GGCGATGGCT TTTATGCGAT GGATTATTGG GGCCAAGGCA CCCTGGTGAC
CCGCTACCGA AAATACGCTA CTAATAAACC CCGGTTCCGT GGGACCACTG

V S S
B1pI

GGTTAGCTCA G
CCAATCGAGT C

```

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Figure 5C: V heavy chain 2 (VH2) gene sequence

```

Q V Q L K E S G P A L V K P T Q T
MfeI
~~~~~
CAGGTGCAAT TGAAGAAG CGGCCCGGCC CTGGTGAAAC CGACCCAAAC
GTCCACGTTA ACTTCTTTC GCCGGGCCGG GACCACTTTG GCTGGGTTTG

L T L T C T F S G F S L S T S G
BspEI
~~~~~
CCTGACCCCTG ACCTGTACCT TTTCGGGATT TAGCCTGTCC ACGTCTGGCG
GGACTGGGAC TGGACATGGA AAAGGCCTAA ATCGGACAGG TGCAGACCCG

V G V G W I R Q P P G K A L E W L
BstXI XhoI
~~~~~
TTGGCGTGGG CTGGATTGCG CAGCCGCCCTG GGAAAGCCCT CGAGTGGCTG
AACCGCACCC GACCTAAGCG GTCGGCGGAC CCTTTCGGGA GCTCACCCGAC

A L I D W D D D K Y Y S T S L K T
MluI
~~~~~
GCTCTGATTG ATTGGGATGA TGATAAGTAT TATAGCACCA GCCTGAAAC
CGAGACTAAC TAACCCCTACT ACTATTCTATA ATATCGTGGT CGGACTTTTG

```

Figure 5C: V heavy chain 2 (VH2) gene sequence (continued)

```

R L T I S K D T S K N Q V V L T
MluI

NspV

GCGTCTGACC ATTAGCAAAG ATACTTCGAA AAATCAGGTG GTGCTGACTA
CGCAGACTGG TAATCGTTTC TATGAAGCTT TTAGTCCAC CACGACTGAT

M T N M D P V D T A T Y Y C A R W
 BSSHII

TGACCAACAT GGACCCGGTG GATACGGCCA CCTATTATTG CGCGCGTTGG
ACTGGTTGTA CCTGGGCCAC CTATGCCGGT GGATAATAAC GCGCGCAACC

G G D G F Y A M D Y W G Q G T L V
 StyI

GGCGGCGATG GCTTTTATGC GATGGATTAT TGGGGCCAAG GCACCCTGGT
CCGCCGCTAC CGAAATATACG CTACCTAATA ACCCCGGTTC CGTGGGACCA

T V S S
 BlnI

GACGGTTAGC TCAG
CTGCCAATCG AGTC

```



Figure 5D: V heavy chain 3 (VH3) gene sequence

```

E V Q L V E S G G G L V Q P G G S
MfeI

GAAGTGCAAT TGGTGAAAG CGGCGGCGGC CTGGTGCAAC CGGGCGGCAG
CTTCACGTTA ACCACCTTTC GCCGCCGCCG GACCACGTTG GCCCGCCGTC

L R L S C A A S G F T F S S Y A
BspEI

CCTGCGTCTG AGCTGCGCGG CCTCCGGATT TACCTTTAGC AGCTATGCCA
GGACGCAGAC TCGACGCGCC GGAGGCCTAA ATGGAATCG TCGATACGCT

M S W V R Q A P G K G L E W V S A
BstXI XhoI

TGAGCTGGGT GCGCCAAGCC CCTGGGAAGG GTCTCGAGTG GGTGAGCGCG
ACTCGACCCA CGCGGTTCGG GGACCCCTCC CAGAGCTCAC CCACTCGCGC

I S G S G G S T Y Y A D S V K G R
ATTAGCGGTA GCGCGGCAG CACCTATTAT GCGGATAGCG TGAAGGCCG
TAATCGCCAT CGCCGCCGTC GTGGATAATA CGCCTATCGC ACTTCCGCGC

```

Figure 5D: V heavy chain 3 (VH3) gene sequence (continued)

```

F T I S R D N S K N T L Y L Q M
 PmlI NspV
      ~~~~~
TTTTACCATT TCACGTGATA ATTCGAAAAA CACCTGTAT CTGCAATGA
AAAATGGTAA AGTGCACTAT TAAGCTTTT GTGGACATA GACGTTACT

N S L R A E D T A V Y C A R W G
      EagI      BssHII
      ~~~~~
ACAGCCTCGG TCGGAAGAT ACGCCCGTGT ATTATTGCGC GCGTTGGGC
TGTCGGACGC ACGCCTTCTA TGCCGGCACA TAATAACGCG CGCAACCCCG

G D G F Y A M D Y W G Q G T L V T
 StyI
      ~~~~~
GGCGATGGCT TTTATGCCGAT GGATTATTGG GGCCAAGGCA CCCTGGTGAC
CCGCTACCGA AAATACGCTA CCTAATAACC CCGGTTCCGT GGGACCACTG

V S S
      BlnI
      ~~~~~
GGTAGCTCA G
CCAATCGAGT C

```

Figure 5E: V heavy chain 4 (VH4) gene sequence

```

Q V Q L Q E S G P G L V K P S E T
 MfeI
~~~~~
CAGGTGCAAT TGCAAGAAAG TGGTCCGGGC CTGGTGAAAC CGAGCGAAAC
GTCCACGTTA ACGTTCTTTC ACCAGGCCCG GACCACTTTG GCTCGCTTTG

L  S  L  T  C  T  V  S  G  G  S  I  S  S  Y  Y
      BspEI
~~~~~
CCTGAGCCTG ACCTGCACCG TTTCGGGAGG CAGCATTAGC AGCTATTATT
GGA CTGGAC TGGACGTGGC AAAGGCCTCC GTCGTAATCG TCGATAATAA

W S W I R Q P P G K G L E W I G Y
 BstXI XhoI
~~~~~
GGAGCTGGAT TCGCCAGCCG CCTGGGAAGG GTCTCGAGTG GATTGGCTAT
CCTCGACCCTA AGCGGTCGGC GGACCCCTTC CAGAGCTCAC CTAACCGATA

I  Y  Y  S  G  S  T  N  Y  N  P  S  L  K  S  R  V
      BstEII
~~~~~
ATTATTATTA GCGGCAGCAC CAACTATAAT CCGAGCCTGA AAAGCCGGGT
TAAATAATAT CGCCGTCGTG GTTGATATTA GGCTCGGACT TTTCGGCCCA

```

Figure 5E: V heavy chain 4 (VH4) gene sequence (continued)

```

 T I S V D T S K N Q F S L K L S
BstEII
~~~~~
GACCATTAGC GTTGATACTT CGAAAACCA GTTAGCCTG AAAC TGAGCA
CTGGTAATCG CAACTATGAA GCTTTTGGT CAAATCGGAC TTTGACTCGT

S V T A A D T A V Y Y C A R W G G
      EagI BssHII
~~~~~
GCGTGACGGC GCGGATACG GCCGTGTATT ATTGCGCGCG TTGGGGCGGC
CGCACTGCCG CCGCCTATGC CGGCACATAA TAACGCGCGC AACCCCGCCG

D G F Y A M D Y W G Q G T L V T V
 StyI
~~~~~
GATGGCTTTT ATGCGATGGA TTATTGGGC CAAGCACCC TGGTGACGGT
CTACCGAAAA TACGCTACCT AATAACCCCG GTTCCGTGGG ACCACTGCCA

```

```

S S
BspI
~~~~~
TAGCTCAG
ATCGAGTC

```

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Figure 5F: V heavy chain 5 (VH5) gene sequence

```

E V Q L V Q S G A E V K K P G E S
MfeI
~~~~~
GAAGTGCAAT TGGTTCAGAG CGGCGCGGAA GTGAAAAAAC CGGCGGAAAG
CTTCACGTTA ACCAAGTCTC GCCGCGCCTT CACTTTTGTG GCCCGCTTTC

L K I S C K G S G Y S F T S Y W
BspEI
~~~~~
CCTGAAAATT AGCTGCAAAG GTTCCGGGATA TTCCTTTACG AGCTATTGGA
GGACTTTTAA TCGACGTTTC CAAGGCCCTAT AAGGAAATGC TCGATAACCT

I G W V R Q M P G K G L E W M G I
BstXI XhoI
~~~~~
TTGGCTGGGT GCGCCAGATG CCTGGGAAGG GTCTCGAGTG GATGGGCATT
AACCGACCCA CGCGGTCTAC GGACCCCTTCC CAGAGCTCAC CTACCCCGTAA

I Y P G D S D T R Y S P S F Q G Q
ATTATCCCG GCGATAGCGA TACCCGTTAT TCTCCGAGCT TTCAGGGCCA
TAAATAGGCC CGCTATCGCT ATGGGCAATA AGAGGCTCGA AAGTCCCGGT

```

Figure 5F: V heavy chain 5 (VH5) gene sequence (continued)

```

V   T   I   S   A   D   K   S   I   S   T   A   Y   L   Q   W
BstEII
~~~~~
GGTGACCATT AGCGGGGATA AAAGCATTAG CACCGCGTAT CTTCAATGGA
CCACTGGTAA TCGCGCCTAT TTTCGTAATC GTGGCGCATA GAAGTTACCT

S S L K A S D T A M Y Y C A R W G
BssHII
~~~~~
GCAGCCTGAA AGCGAGCGAT ACGGCCATGT ATTATTGCGC GCGTTGGGGC
CGTCGGACTT TCGCTCGCTA TGCCGGTACA TAATAACGCG CGCAACCCCG

G   D   G   F   Y   A   M   D   Y   W   G   Q   G   T   L   V   T
StyI
~~~~~
GGCGATGGCT TTTATGCGAT GGATTATTGG GGCCAAGGCA CCCTGGTGAC
CCGCTACCGA AAATACGCTA CCTAATAACC CCGGTTCCGT GGGACCACTG

V S S
BlpI
~~~~~
GGTAGCTCA G
CCAATCGAGT C

```

Figure 5G: V heavy chain 6 (VH6) gene sequence

```

Q V Q L Q Q S G P G L V K P S Q T
MfeI
~~~~~
CAGGTGCAAT TGCAACAGTC TGTCCGGGC CTGGTGAAC CGAGCCAAAC
GTCCACGTTA ACGTTGTCAG ACCAGGCCCG GACCACTTTG GCTCGGTTTG

L S L T C A I S G D S V S S N S
BspEI
~~~~~
CCTGAGCCTG ACCTGTGCGA TTTCCGGAGA TAGCGTGAGC AGCAACAGCG
GGACTCGGAC TGGACACGCT AAAGGCCTCT ATCGCACTCG TCGTTGTCCG

A A W N W I R Q S P G R G L E W L
BstXI XhoI
~~~~~
CGGCGTGGAA CTGGATTTCG CAGTCTCCTG GCGGTGGCCT CGAGTGGCTG
GCCGCACCTT GACCTAAGCG GTCAGAGGAC CCGCACCGGA GTCACCGAC

G R T Y Y R S K W Y N D Y A V S V
GGCCGTACCT ATTATCGTAG CAAATGGTAT AACGATTATG CGGTGAGCGT
CCGGCATGGA TAATAGCATC GTTTACCATA TTGCTAATAC GCCACTCGCA

```

Figure 5G: V heavy chain 6 (VH6) gene sequence (continued)

```

K S R I T I N P D T S K N Q F S
 BsaBI NspV
      ~~~~~
GAAAAGCCGG ATTACCATCA ACCCGGATAC TTCGAAAAC CAGTTAGCC
CTTTTCGGCC TAATGGTAGT TGGGCCTATG AAGCTTTTG GTCAAATCGG

L Q L N S V T P E D T A V Y C A
      EagI      BssHII
      ~~~~~
TGCAACTGAA CAGCGTGACC CCGGAAGATA CGGCCGTGTA TTATTGCGCG
ACGTTGACTT GTCGCACTGG GGCCTTCTAT GCCGGCACAT AATAACGCGC

R W G G D G F Y A M D Y W G Q G T
 BssHII StyI
      ~~~~~
CGTTGGGGCG GCGATGGCTT TTATGCGATG GATTATTGG GCCAAGGCAC
GCAACCCCGC CGTACCGAA AATACGCTAC CTAATAACCC CGGTCCCGTG

L V T V S S
      BlnI
      ~~~~~
CCTGGTGACG GTTAGCTCAG
GGACCACTGC CAATCGAGTC

```



Figure 6: oligonucleotides for gene synthesis

**O1K1** 5' - GAATGCATACGCTGATATCCAGATGACCCAGAG-  
CCCGTCTAGCCTGAGC -3'

**O1K2** 5' - CGCTCTGCAGGTAATGGTCACACGATCACCCAC-  
GCTCGCGCTCAGGCTAGACGGGC -3'

**O1K3** 5' - GACCATTACCTGCAGAGCGAGCCAGGGCATTAG-  
CAGCTATCTGGCGTGGTACCAGCAG -3'

**O1K4** 5' - CTTTGCAAGCTGCTGGCTGCATAAATTAATAGT-  
TTCGGTGCTTTACCTGGTTTCTGCTGGTACCACGCCAG -3'

**O1K5** 5' - CAGCCAGCAGCTTGCAAAGCGGGGTCCCGTCCC-  
GTTTTAGCGGCTCTGGATCCGGCACTGATTTTAC -3'

**O1K6** 5' - GATAATAGGTCGCAAAGTCTTCAGGTTGCAGGC-  
TGCTAATGGTCAGGGTAAAATCAGTGCCGGATCC -3'

**O2K1** 5' - CGATATCGTGATGACCCAGAGCCCCTGAGCCT-  
GCCAGTGACTCCGGGCGAGCC -3'

**O2K2** 5' - GCCGTTGCTATGCAGCAGGCTTTGGCTGCTTCT-  
GCAGCTAATGCTCGCAGGCTCGCCCGGAGTCAC -3'

**O2K3** 5' - CTGCTGCATAGCAACGGCTATAACTATCTGGAT-  
TGGTACCTTCAAAAACCAGGTCAAAGCCC -3'

**O2K4** 5' - CGATCCGGGACCCCACTGGCACGGTTGCTGCCC-  
AGATAAATTAATAGCTGCGGGCTTTGACCTGGTTTTTG -3'

**O2K5** 5' - AGTGGGGTCCCGGATCGTTTTAGCGGCTCTGGA-  
TCCGGCACCGATTTTACCCTGAAAATTAGCCGTGTG -3'

**O2K6** 5' - CCATGCAATAATACACGCCACGTCTTCAGCTT-  
CCACACGGCTAATTTTCAGGG -3'

**O3K1** 5' - GAATGCATACGCTGATATCGTGCTGACCCAGAG-  
CCCGG -3'

**O3K2** 5' - CGCTCTGCAGCTCAGGGTCGCACGTTGCCCCG-  
AGACAGGCTCAGGGTCGCCGGGCTCTGGGTCAGC -3'

**O3K3** 5' - CCCTGAGCTGCAGAGCGAGCCAGAGCGTGAGCA-  
GCAGCTATCTGGCGTGGTACCAG -3'

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Figure 6: (continued)

**O3K4** 5' - GCACGGCTGCTCGCGCCATAAATTAATAGACGC-  
GGTGCTTGACCTGGTTTCTGCTGGTACCACGCCAGATAG -3'

**O3K5** 5' - GCGCGAGCAGCCGTGCAACTGGGGTCCCGGCGC-  
GTTTTAGCGGCTCTGGATCCGGCACGGATTTTAC -3'

**O3K6** 5' - GATAATACACCGCAAAGTCTTCAGGTTCCAGGC-  
TGCTAATGGTCAGGGTAAAATCCGTGCCGGATC -3'

**O4K1** 5' - GAATGCATACGCTGATATCGTGATGACCCAGAG-  
CCCGGATAGCCTGGCG -3'

**O4K2** 5' - GCTTCTGCAGTTAATGGTCGCACGTTCGCCCAG-  
GCTCACCGCCAGGCTATCCGGGC -3'

**O4K3** 5' - CGACCATTAAGTGCAGAAGCAGCCAGAGCGTGC-  
TGTATAGCAGCAACAACAAAACCTATCTGGCGTGGTACCAG -  
3'

**O4K4** 5' - GATGCCCAATAAATTAATAGTTTCGGCGGCTGA-  
CCTGGTTTCTGCTGGTACCACGCCAGATAG -3'

**O4K5** 5' - AAATATTAATTTATTGGGCATCCACCCGTGAA-  
AGCGGGGTCCCGGATCGTTTTAGCGGCTCTGGATCCGGCAC-  
3'

**O4K6** 5' - GATAATACACCGCCACGTCTTCAGCTTGCAGGG-  
ACGAAATGGTCAGGGTAAAATCAGTGCCGGATCCAGAGCC -  
3'

**O1L1** 5' - GAATGCATACGCTCAGAGCGTGCTGACCCAGCC-  
GCCTTCAGTGAGTGG -3'

**O1L2** 5' - CAATGTTGCTGCTGCTGCCGCTACACGAGATGG-  
TCACACGCTGACCTGGTGCGCCACTCACTGAAGGCGGC -3'

**O1L3** 5' - GGCAGCAGCAGCAACATTGGCAGCAACTATGTG-  
AGCTGGTACCAGCAGTTGCCCCGGGAC -3'

**O1L4** 5' - CCGGCACGCCTGAGGGACGCTGGTTGTTATCAT-  
AAATCAGCAGTTTCGGCGCCGTCCCGGGCAACTGC -3'

**O1L5** 5' - CCCTCAGGCGTGCCGGATCGTTTTAGCGGATCC-  
AAAAGCGGCACCAGCGCGAGCCTTGCG -3'

Figure 6: (continued)

**O1L6** 5' - CCGCTTCGTCTTCGCTTTGCAGGCCCGTAATCG-  
CAAGGCTCGCGCTGG -3'

**O2L1** 5' - GAATGCATACGCTCAGAGCGCACTGACCCAGCC-  
AGCTTCAGTGAGCGGC -3'

**O2L2** 5' - CGCTGCTAGTACCCGTACACGAGATGGTAATGC-  
TCTGACCTGGTGAGCCGCTCACTGAAGCTGG -3'

**O2L3** 5' - GTACGGGTACTAGCAGCGATGTGGGCGGCTATA-  
ACTATGTGAGCTGGTACCAGCAGCATCCCGG -3'

**O2L4** 5' - CGCCTGAGGGACGGTTGCTCACATCATAAATCA-  
TCAGTTTCGGCGCCTTCCCGGGATGCTGCTGGTAC -3'

**O2L5** 5' - CAACCGTCCCTCAGGCGTGAGCAACCGTTTTAG-  
CGGATCCAAAAGCGGCAACACCGCGAGCC -3'

**O2L6** 5' - CCGCTTCGTCTTCCGCTTGCAGGCCGCTAATGG-  
TCAGGCTCGCGGTGTTGCCG -3'

**O3L1** 5' - GAATGCATACGCTAGCTATGAACTGACCCAGCC-  
GCCTTCAGTGAGCG -3'

**O3L2** 5' - CGCCCAGCGCATCGCCGCTACACGAGATACGCG-  
CGGTCTGACCTGGTGCAACGCTCACTGAAGGCGGC -3'

**O3L3** 5' - GGCGATGCGCTGGGCGATAAATACGCGAGCTGG-  
TACCAGCAGAAACCCGGGCAGGCGC -3'

**O3L4** 5' - GCGTTCCGGGATGCCTGAGGGACGGTCAGAATC-  
ATCATAAATCACCAGAACTGGCGCCTGCCCCGGGTTC -3'

**O3L5** 5' - CAGGCATCCCGGAACGCTTTAGCGGATCCAACA-  
GCGGCAACACCGCGACCCTGACCATTAGCGG -3'

**O3L6** 5' - CCGCTTCGTCTTCCGCCTGAGTGCCGCTAATGG-  
TCAGGGTC -3'

**O1246H1** 5' - GCTCTTCACCCCTGTTACCAAAGCCCAG-  
GTGCAATTG -3'

**O1AH2** 5' - GGCTTTGCAGCTCACTTTCACGCTGCTGCCCCGG-  
TTTTTTCACCTCCGCGCCAGACTGAACCAATTGCACCTGGGC-  
TTTG -3'

Figure 6: (continued)

**01AH3** 5' - GAAAGTGAGCTGCAAAGCCTCCGGAGGCACTTT-  
TAGCAGCTATGCGATTAGCTGGGTGCGCCAAGCCCCTGGGCAG  
GGTC -3'

**01AH4** 5' - GCCCTGAAACTTCTGCGCGTAGTTCGCCGTGCC-  
AAAAATCGGAATAATGCCGCCCATCCACTCGAGACCCTGCCC-  
AGGGGC -3'

**01AH5** 5' - GCGCAGAAGTTTCAGGGCCGGGTGACCATTACC-  
GCGGATGAAAGCACCAGCACC GCGTATATGGA ACTGAGCAGCC  
TGCG -3'

**01ABH6** 5' - GCGCGCAATAATACAGGCCGTATCTTCGCT-  
ACGCAGGCTGCTCAGTTCC -3'

**01BH2** 5' - GGCTTTGCAGCTCACTTTCACGCTCGCGCCCGG-  
TTTTTTC ACTTCCGCGCCGCTCTGAACCAATTGCACCTGGGC-  
TTTG -3'

**01BH3** 5' - GAAAGTGAGCTGCAAAGCCTCCGGATATACCTT-  
TACCAGCTATTATATGCACTGGGTCCGCCAAGCCCCTGGGCAG  
GGTC -3'

**01BH4** 5' - GCCCTGAAACTTCTGCGCGTAGTTCGTGCCGCC-  
GCTATTCGGGTTAATCCAGCCCATCCACTCGAGACCCTGCCCCA  
GGGGC -3'

**01BH5** 5' - GCGCAGAAGTTTCAGGGCCGGGTGACCATGACC-  
CGTGATAACCAGCATTAGCACC GCGTATATGGA ACTGAGCAGCC  
TGCG -3'

**02H2** 5' - GGTACAGGTCAGGGTCAGGGTTTGGGTCCGTTT-  
CACCAGGGCCGGGCCGCTTTCTTTCAATTGCACCTGGGCTTTG  
-3'

**02H3** 5' - CTGACCCTGACCTGTACCTTTTCCGGATTTAGC-  
CTGTCCACGTCTGGCGTTGGCGTGGGCTGGATTCGCCAGCCGC  
CTGGGAAAG -3'

**02H4** 5' - GCGTTTTTCAGGCTGGTGCTATAATACTTATCAT-  
CATCCCAATCAATCAGAGCCAGCCACTCGAGGGCTTTCCCAGG  
CGGCTGG -3'

Figure 6: (continued)

**O2H5** 5' - GCACCAGCCTGAAAACGCGTCTGACCATTAGCA-  
AAGATACTTCGAAAAATCAGGTGGTGCTGACTATGACCAACAT  
GG -3'

**O2H6** 5' - GCGCGCAATAATAGGTGGCCGTATCCACCGGGT-  
CCATGTTGGTCATAGTCAGC -3'

**O3H1** 5' - CGAAGTGCAATTGGTGGAAAGCGGCGGCGGCCT-  
GGTGCAACCGGGCGGCAG -3'

**O3H2** 5' - CATAGCTGCTAAAGGTAAATCCGGAGGCCGCGC-  
AGCTCAGACGCAGGCTGCCGCCCGGTTGCAC -3'

**O3H3** 5' - GATTTACCTTTAGCAGCTATGCGATGAGCTGGG-  
TGCGCCAAGCCCCCTGGGAAGGGTCTCGAGTGGGTGAG -3'

**O3H4** 5' - GGCCTTTCACGCTATCCGCATAATAGGTGCTGC-  
CGCCGCTACCGCTAATCGCGCTCACCCACTCGAGACCC -3'

**O3H5** 5' - CGGATAGCGTGAAAGGCCGTTTTACCATTTCAC-  
GTGATAATTTCGAAAAACACCCTGTATCTGCAAATGAACAG-3'

**O3H6** 5' - CACGCGCGCAATAATACACGGCCGTATCTTCCG-  
CACGCAGGCTGTTTCATTTGCAGATACAGG -3'

**O4H2** 5' - GGTCAGGCTCAGGGTTTCGCTCGGTTTCACCAG-  
GCCCCGGACCACTTTCTTGCAATTGCACCTGGGCTTTG -3'

**O4H3** 5' - GAAACCCTGAGCCTGACCTGCACCGTTTCCGGA-  
GGCAGCATTAGCAGCTATTATTGGAGCTGGATTCGCCAGCCGC  
-3'

**O4H4** 5' - GATTATAGTTGGTGCTGCCGCTATAATAAATAT-  
AGCCAATCCACTCGAGACCCTTCCCAGGCGGCTGGCGAATCCA  
G -3'

**O4H5** 5' - CGGCAGCACCAACTATAATCCGAGCCTGAAAAG-  
CCGGGTGACCATTAGCGTTGATACTTCGAAAAACCAGTTTAGC  
CTG -3'

**O4H6** 5' - GCGCGCAATAATACACGGCCGTATCCGCCGCCG-  
TCACGCTGCTCAGTTTCAGGCTAAACTGGTTTTTCG -3'

Figure 6: (continued)

**05H1** 5' - GCTCTTCACCCCTGTTACCAAAGCCGAAGTGCA-  
ATTG -3'

**05H2** 5' - CCTTTGCAGCTAATTTTCAGGCTTTCGCCCCGGT-  
TTTTTCACTTCCGCGCCGCTCTGAACCAATTGCACTTCGGCTT  
TGG -3'

**05H3** 5' - CCTGAAAATTAGCTGCAAAGGTTCCGGATATTC-  
CTTTACGAGCTATTGGATTGGCTGGGTGCGCCAGATGCCTGG  
-3'

**05H4** 5' - CGGAGAATAACGGGTATCGCTATCGCCCCGGATA-  
AATAATGCCCATCCACTCGAGACCCTTCCCAGGCATCTGGCGC  
AC -3'

**05H5** 5' - CGATACCCGTTATTCTCCGAGCTTTCAGGGCCA-  
GGTGACCATTAGCGCGGATAAAAGCATTAGCACCGCGTATCTT  
C -3'

**05H6** 5' - GCGCGCAATAATACATGGCCGTATCGCTCGCTT-  
TCAGGCTGCTCCATTGAAGATACGCGGTGCTAATG -3'

**06H2** 5' - GAAATCGCACAGGTCAGGCTCAGGGTTTGGCTC-  
GGTTTCACCAGGCCCCGGACCAGACTGTTGCAATTGCACCTGG-  
GCTTTG -3'

**06H3** 5' - GCCTGACCTGTGCGATTTCCGGAGATAGCGTGA-  
GCAGCAACAGCGCGGCGTGGAAGTTCGCCAGTCTCCTGG  
GCG -3'

**06H4** 5' - CACCGCATAATCGTTATACCATTTGCTACGATA-  
ATAGGTACGGCCCAGCCACTCGAGGCCACGCCCAGGAGACTG-  
GCG -3'

**06H5** 5' - GGTATAACGATTATGCGGTGAGCGTGAAAAGCC-  
GGATTACCATCAACCCGGATACTTCGAAAAACCAGTTTAGCCT  
GC -3'

**06H6** 5' - GCGCGCAATAATACACGGCCGTATCTTCCGGGG-  
TCACGCTGTTCAAGTTGCAGGCTAAACTGGTTTTTC -3'

**OCLK1** 5' - GGCTGAAGACGTGGGCGTGTATTATTGCCAGCA-  
GCATTATACCACCCCGCCGACCTTTGGCCAGGGTAC -3'

Figure 6: (continued)

OCLK2 5' - GCGGAAAAATAAACACGCTCGGAGCAGCCACCG-  
TACGTTTAATTTCAACTTTCGTACCCTGGCCAAAGGTC -3'

OCLK3 5' - GAGCGTGTTTATTTTTCCGCCGAGCGATGAACA-  
ACTGAAAAGCGGCACGGCGAGCGTGGTGTGCCTGCTG -3'

OCLK4 5' - CAGCGCGTTGTCTACTTTCCACTGAACTTTCGC-  
TTCACGCGGATAAAAGTTGTTTCAGCAGGCACACCACGC -3'

OCLK5 5' - GAAAGTAGACAACGCGCTGCAAAGCGGCAACAG-  
CCAGGAAAGCGTGACCGAACAGGATAGCAAAGATAG -3'

OCLK6 5' - GTTTTTTCATAATCCGCTTTGCTCAGGGTCAGGG-  
TGCTGCTCAGAGAATAGGTGCTATCTTTGCTATCCTGTTTCG -  
3'

OCLK7 5' - GCAAAGCGGATTATGAAAAACATAAAGTGTATG-  
CGTGCGAAGTGACCCATCAAGGTCTGAGCAGCCCCGGTG -3'

OCLK8 5' - GGCATGCTTATCAGGCCTCGCCACGATTAAAAG-  
ATTTAGTCACCGGGCTGCTCAGAC -3'

OCH1 5' - GGCGTCTAGAGGCCAAGGCACCCTGGTGACGGT-  
TAGCTCAGCGTCGAC -3'

OCH2 5' - GTGCTTTTGCTGCTCGGAGCCAGCGGAAACACG-  
CTTGGACCTTTGGTCGACGCTGAGCTAACC -3'

OCH3 5' - CTCCGAGCAGCAAAAGCACCAGCGGCGGCACGG-  
CTGCCCTGGGCTGCCTGGTTAAAGATTATTTCC -3'

OCH4 5' - CTGGTCAGCGCCCCGCTGTTCCAGCTCACGGTG-  
ACTGGTTCCGGGAAATAATCTTTAACCAGGCA -3'

OCH5 5' - AGCGGGGCGCTGACCAGCGGCGTGACATACCTTT-  
CCGGCGGTGCTGCAAAGCAGCGGCCTG -3'

OCH6 5' - GTGCCTAAGCTGCTGCTCGGCACGGTCACAACG-  
CTGCTCAGGCTATACAGGCCGCTGCTTTGCAG -3'

OCH7 5' - GAGCAGCAGCTTAGGCACTCAGACCTATATTTG-  
CAACGTGAACCATAAACCGAGCAACACC -3'

OCH8 5' - GCGCGAATTCGCTTTTCGGTTCCACTTTTTTAT-  
CCACTTTGGTGTGCTCGGTTTATGG -3'

Figure 7A: sequence of the synthetic Cx gene segment

```

 ° V A A A P S V F I F P P S D E Q
 BsiWI

CGTACGGTGG CTGCTCCGAG CGTGTTTATT TTTCCGCCGA GCGATGAACA
GCATGCCACC GACGAGGCTC GCACAAATAA AAAGCGGCT CGTACTTGT

 L K S G T A S V V C L L N N F Y
ACTGAAAAGC GGCACGGCGA GCGTGGTGTG CCTGCTGAAC AACTTTTATC
TGACTTTTCG CCGTGCCGCT CGCACCCACAC GGACGACTTG TTGAAAATAG

 P R E A K V Q W K V D N A L Q S G
CGCGTGAAGC GAAAGTTCAG TGGAAAGTAG ACAACGCGCT GCAAAGCGGC
GGCACTTCG CTTTCAAGTC ACCTTTCATC TGTTGCGCGA CGTTTCGCCG

 N S Q E S V T E Q D S K D S T Y S
AACAGCCAGG AAAGCGTGAC CGAACAGGAT AGCAAAGATA GCACCTATTC
TTGTCGGTCC TTTCCGCACTG GCTGTCCCTA TCGTTTCTAT CGTGGATAAG

 L S S T L T L S K A D Y E K H K
TCTGAGCAGC ACCCTGACCC TGAGCAAAGC GGATTATGAA AAACATAAAG
AGACTCGTCG TGGGACTGGG ACTCGTTTCG CTAATACTT TTTGTATTTC

```



Figure 7A: sequence of the synthetic Cx gene segment (continued)

V Y A C E V T H Q G L S S P V T K  
 TGTATGCGTG CGAAGTGACC CATCAAGGTC TGAGCAGCCCC GGTGACTAAA  
 ACATACGCAC GCTTCACTGG GTAGTCCAG ACTCGTCGGG CCACTGATTT

S F N R G E A \*  
 StuI SphI  
 ~~~~~  
 TCCTTTAATC GTGGCGAGGC CTGATAAGCA TGC  
 AGAAAATTAG CACCGCTCCG GACTATTCGT ACG

Figure 7B: sequence of the synthetic CH1 gene segment

```

A S T K G P S V F P L A P S S
BlpI Sali
~~~~~
GCTCAGCGTC GACCAAAGGT CCAAGCGTGT TTCCGCTGGC TCCGAGCAGC
CGAGTCGCAG CTGGTTTCCA GGTTCGCACA AAGGCGACCG AGGCTCGTCG

K S T S G G T A A L G C L V K D Y
AAAAGCACCA GCGCGGCAC GGCTGCCCTG GGTGCCCTGG TTAAGATTA
TTTTCGTGGT CGCGCCCGTG CCGACGGGAC CCGACGGACC AATTCTAAT

F P E P V T V S W N S G A L T S
TTTCCCGGAA CCAGTCACCG TGAGCTGGAA CAGCGGGCGG CTGACCAGCG
AAAGGGCCTT GGTCA GTGC ACTCGACCTT GTCGCCCGC GACTGGTCGC

G V H T F P A V L Q S S G L Y S L
GCGTGCATAC CTTCCGGCG GTGCTGCAAA GCAGCGGCCT GTATAGCCTG
CGCACGTATG GAAAGGCCG CACGACGTTT CGTCGCCGGA CATATCGGAC

S S V V T V P S S S L G T Q T Y I
AGCAGCGTTG TGACCGTGCC GAGCAGCAGC TTAGGCACTC AGACCTATAT
TCGTCGCAAC ACTGGCACGG CTCGTCTGTCG AATCCGTGAG TCTGGATATA

```

Figure 7B: sequence of the synthetic CH1 gene segment (continued)

C	N	V	N	H	K	P	S	N	T	K	V	D	K	K	V
TTGCAACGTG	AACCATAAAC	CGAGCAACAC	CAAAGTGGAT	AAAAAAGTGG											
AACGTTGCAC	TTGGTATTG	GTCGTTGTG	GTTTCACCTA	TTTTTTCACC											

E	P	K	S	E	F	*
				EcoRI		HindIII
				~~~~~		~~~~~
AACCGAAAAG	CGAATTCTGA	TAAGCTT				
TTGGCTTTC	GCTTAAGACT	ATTCGAA				

Figure 7C: functional map and sequence of module 24 comprising the synthetic Cλ gene segment (huCL lambda)

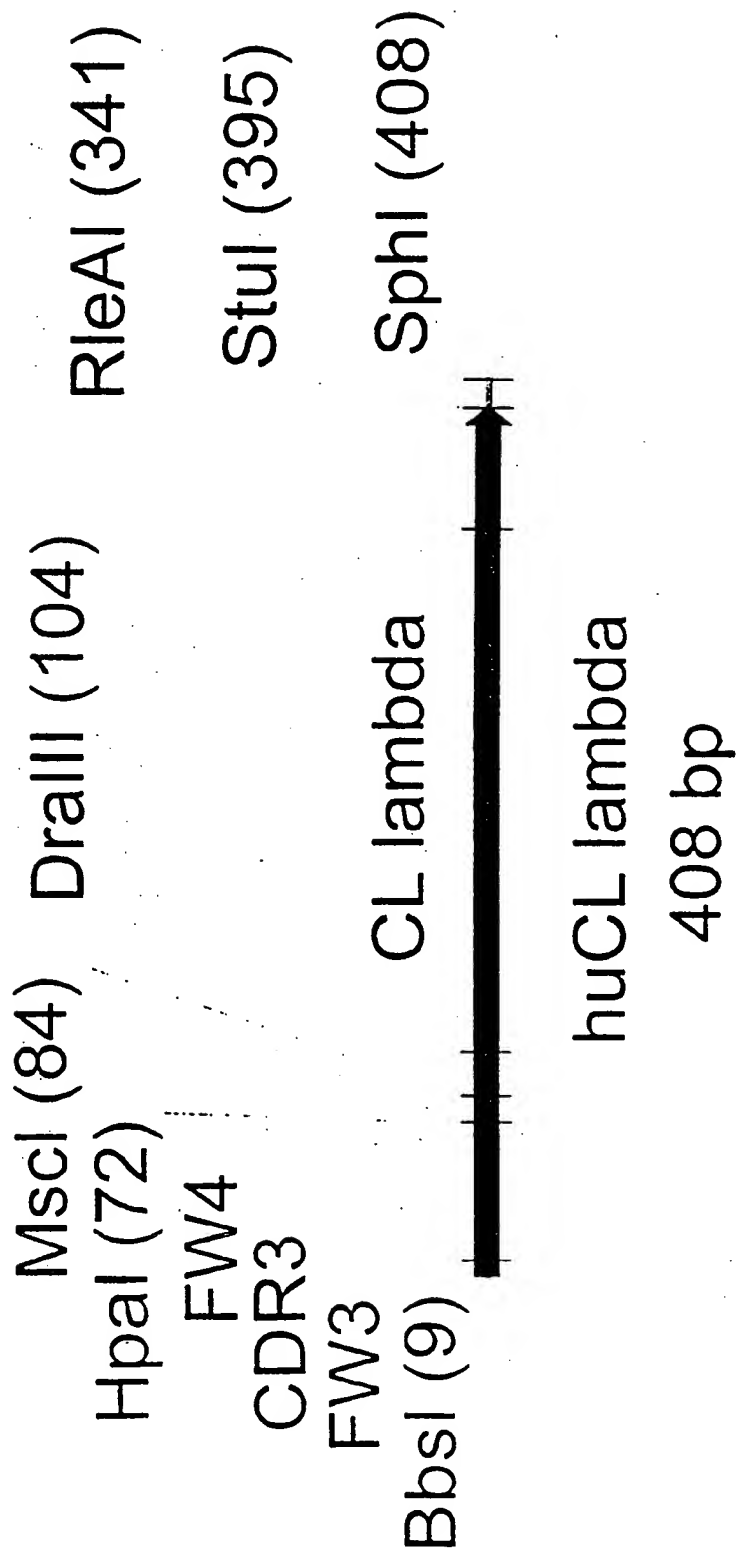


Figure 7C: functional map and sequence of module 24 comprising the synthetic Cl gene segment (huCL lambda) (continued)

	Bbs I		Hpa I	Msc I	Dra III
	~~~~~		~~~~~	~~~~~	~~~~~
1	GAAGACGAAG	CGGATTATTA	TTGCCAGCAG	CATTATACCA	CCCCGCCTGT
	CTTCTGCTTC	GCCTAATAAT	AACGGTCGTC	GTAATATGGT	GGGGCGGACA
51	GTTTGGCGC	GGCACGAAGT	TAACCGTTCT	TGGCCAGCCG	AAAGCCGCAC
	CAAACCGCCG	CCGTGCTTCA	ATTGGCAAGA	ACCGGTCGGC	TTTCGGCGTG
	Dra III				
	~~~~~				
101	CGAGTGTGAC	GCTGTTTCCG	CCGAGCAGCG	AAGAATTGCA	GGCGAACAAA
	GCTCACACTG	CGACAAAGGC	GGCTCGTCGC	TTCTTAACGT	CCGCTTGTTT
151	GCGACCCCTGG	TGTGCCTGAT	TAGCGACTTT	TATCCGGGAG	CCGTGACAGT
	CGCTGGGACC	ACACGACTA	ATCGCTGAAA	ATAGGCCCTC	GGCACTGTCA
201	GGCCTGGAAG	GCAGATAGCA	GCCCCGTCAA	GGCGGGAGTG	GAGACCACCA
	CCGGACCTTC	CGTCTATCGT	CGGGGCAGTT	CCGCCCTCAC	CTCTGGTGGT

Figure 7C: functional map and sequence of module 24 comprising the synthetic CI gene segment (huCl lambda) (continued)

251 CACCCTCCAA ACAAGCAAC AACAGTACG CGGCCAGCAG CTATCTGAGC
GTGGGAGGTT TGTTTCGTTG TTGTTTCATGC GCCGGTCGTC GATAGACTCG

RleAI

~~~~~

301 CTGACGCCCTG AGCAGTGGAA GTCCACACAGA AGCTACAGCT GCCAGGTCAC  
GACTGCGGGAC TCGTCACCTT CAGGGTGTCT TCGATGTCGA CGGTCCAGTG

StuI

~~~~~

351 GCATGAGGGG AGCACC GTGG AAAAACC GT TGC GCCGACT GAGGCCTGAT
CGTACTCCCC TCGTGGCACC TTTT TGGCA ACGCGGCTGA CTCCGGACTA

SphI

~~~~~

401 AAGCATGC  
TTCGTACG

Figure 7D: oligonucleotides used for synthesis of module M24 containing Cλ gene segment

M24: assembly PCR

M24-A: GAAGACAAGCGGATTATTATTGCCAGCAGCATTATACCAACCCGCCCTGTGTTGGCGGCG-

GCACGAAGTTAACCGTTC

M24-B: CAATTCTCGCTGCTCGGCGGAACAGCGTCACACTCGGTGCGGCTTCGGCTGGCCAA-

GAACGGTTAACTTCGTGCCGC

M24-C: CGCCGAGCAGCGAAGAATTGCAGGCGAACAAGCGACCCCTGGTGTGCCCTGATTAGCGACT-

TTTATCCGGGAGCCGTGACA

M24-D: TGTTGGAGGGTGTGGTGTCTCCACTCCCGCCTTGACGGGGCTGCTAICTGCCCTCCAG-

GCCACTGTCACGGCTCCCGG

M24-E: CCACACCTCCAAACAAGCAACAAGTACGCGGCCAGCAGCTATCTGAGCCTGACGC-

CTGAGCAGTGGAAGTCCACAGAAGCTACAGCTG

M24-F: GCATGCTTATCAGGCCTCAGTCGGCGCAACGGTTTTTCCACGGTGTCCCTCATGCCGT-

GACCTGGCAGCTGTAGCTTC

Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2

|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |             |            |             |             |       |   |   |   |   |   |   |   |   |   |   |   |
|------------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|-------------|------------|-------------|-------------|-------|---|---|---|---|---|---|---|---|---|---|---|
| M          | K | Q | S | T | I | A | L | A | L | L | P | L | L | F | T | P           |            |             |             |       |   |   |   |   |   |   |   |   |   |   |   |
|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | SapI        |            |             |             |       |   |   |   |   |   |   |   |   |   |   |   |
|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | -----       |            |             |             |       |   |   |   |   |   |   |   |   |   |   |   |
| ATGAAACAAA |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | GCACTATTGC  | ACTGGCACTC | TTACCGTTGC  | TCTTCACCCC  |       |   |   |   |   |   |   |   |   |   |   |   |
| TACTTTGTTT |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | CGTGATAACG  | TGACCGTGAG | AATGGCAACG  | AGAAGTGGGG  |       |   |   |   |   |   |   |   |   |   |   |   |
| V          |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | T           | K          | A           | D           | Y     | K | D | E | V | Q | L | V | E | S | G |   |
|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |             |            |             |             | MfeI  |   |   |   |   |   |   |   |   |   |   |   |
|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |             |            |             |             | ----- |   |   |   |   |   |   |   |   |   |   |   |
| TGTTACCAAA |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | GCCGACTACA  | AAGATGAAGT | GCAATTGGTG  | GAAAGCGGCG  |       |   |   |   |   |   |   |   |   |   |   |   |
| ACAATGGTTT |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | CGGCTGATGT  | TTCTACTTCA | CGTTAACCCAC | CTTTCGCCCG  |       |   |   |   |   |   |   |   |   |   |   |   |
| G          |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | G           | L          | V           | Q           | P     | G | G | S | L | R | L | S | C | A | A | S |
|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |             |            |             |             | BspEI |   |   |   |   |   |   |   |   |   |   |   |
|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |             |            |             |             | ----- |   |   |   |   |   |   |   |   |   |   |   |
| GCGGCCTGGT |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | GCAACCGGGC  | GGCAGCCTGC | GTCTGAGCTG  | CGCGGCCTCC  |       |   |   |   |   |   |   |   |   |   |   |   |
| CGCCGGACCA |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | CGTTGGCCCCG | CCGTCGGACG | CAGACTCGAC  | GCGCCGGAGG  |       |   |   |   |   |   |   |   |   |   |   |   |
| G          |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | F           | T          | F           | S           | S     | Y | A | M | S | W | V | R | Q | A | P | G |
|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |             |            |             |             | BspEI |   |   |   |   |   |   |   |   |   |   |   |
|            |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |             |            |             |             | ----- |   |   |   |   |   |   |   |   |   |   |   |
| GGATTACCT  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | TTAGCAGCTA  | TGCGATGAGC | TGGGTGCGCC  | AAGCCCCCTGG |       |   |   |   |   |   |   |   |   |   |   |   |
| CCTAAATGGA |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | AATCGTCGAT  | ACGCTACTCG | ACCCACGCGG  | TTCGGGGACC  |       |   |   |   |   |   |   |   |   |   |   |   |



Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2 (continued)

```

      K  G  L  E  W  V  S  A  I  S  G  S  G  S  T
      XhoI
      -----
GAAGGGTCTC GAGTGGGTGA GCGCGATTAG CCGTAGCGGC GGCAGCACCT
CTTCCCAGAG CTCACCCACT CCGGCTAATC GCCATCGCCG CCGTCGTGGA

Y  Y  A  D  S  V  K  G  R  F  T  I  S  R  D  N  S
      PmlI      NspV
      -----
ATTATGCGGA TAGCGTGAAA GCGCGTTTTC CCATTTCACG TGATAATTTCG
TAATACGCCT ATCGCACTTT CCGGCAAAAT GGTAAGTGC ACTATTAAGC

K  N  T  L  Y  L  Q  M  N  S  L  R  A  E  D  T  A
      NspV      EagI
      -----
AAAAACACCC TGTATCTGCA AATGAACAGC CTGCGTGCGG AAGATACGGC
TTTTTGTTGG ACATAGACGT TTAAGTGTGC GACGCACGCC TTCTATGCCG

V  Y  Y  C  A  R  W  G  G  D  G  F  Y  A  M  D
      EagI      BssHII
      -----
CGTGTATTAT TCGCGCGGTT GGGCGGCGA TGGCTTTTAT GCGATGGATT

```

Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2 (continued)

```

GCACATAATA ACGCGCGCAA CCCGCGGCT ACCGAAATA CGCTACCTAA
Y W G Q G T L V T V S S A G G G S
                               B1pI
                               -----
ATTGGGGCCA AGCACCCCTG GTGACGGTTA GCTCAGCGGG TGGCGGTTCT
TAACCCCGGT TCCGTGGGAC CACTGCCAAT CGAGTCGCCC ACCGCCAAGA

G G G G S G G G G G S G G G S D I
                                EcorV
                                -----
GGCGGCGGTG GGAGCGGTGG CCGTGTTCTT GCGGTGGTG GTTCCGATAT
CCGCGGCCAC CCTCGCCACC GCCACCAAGA CCGCCACCAC CAAGGCTATA

V M T Q S P L S L P V T P G E P
EcorV      BanII
-----
CGTGATGACC CAGAGCCAC TGAGCCTGCC AGTACTCCG GCGAGCCTG
GCACTACTGG GTCTCGGGTG ACTCGGACGG TCACTGAGGC CCGCTCGGAC

A S I S C R S S Q S L L H S N G Y
                               PstI
                               -----
CGAGCATTAG CTGCAGAAGC AGCCAAAGCC TGCTGCATAG CAACGGCTAT
GCTCGTAATC GACGTCTTCG TCGGTTTCGG ACGACGTATC GTTGCCGATA

```

Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2 (continued)

```

N Y L D W Y L Q K P G Q S P Q L L
kpnI      SexAI      AseI
-----
AACTATCTGG ATGGGTACCT TC AAAA ACCA GTCAAAGCC CGCAGCTATT
TTGATAGACC TAACCATGGA AGTTT TGGT CCAGTTTCGG GCGTCGATAA

I Y L G S N R A S G V P D R F S
AseI      Eco109I
-----
AATTATCTG GGCAGCAACC GTGCCAGTGG GTCCC GGAT CGTTTAGCG
TTAAATAGAC CCGTCGTTGG CACGGTCACC CCAGGGCCTA GCAAAATCGC

G S G S G T D F T L K I S R V E A
BamHI
-----
GCTCTGGATC CGGCACCGAT TTACCCCTGA AAATTAGCCG TGTGGAAGCT
CGAGACCTAG GCCGTGGCTA AAATGGGACT TTAATCGGC ACACCTTCGA

E D V G V Y Y C Q Q H Y T P P T
BbsI
-----
GAAGACGTGG GCGTGATTA TTGCCAGCAG CATTATACCA CCCC GCCGAC
CTTCTGCACC CGCACATAAT AACGGTCGTC GTAATATGGT GGGCGGCTG

```

Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2 (continued)

|                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |
|--------------------------------------------------|---|---|---|---|---|---|---|---|---|---|---|---|---|
| F                                                | G | Q | G | T | K | V | E | I | K | R | T | E | F |
| MSCI                                             |   |   |   |   |   |   |   |   |   |   |   |   |   |
| -----                                            |   |   |   |   |   |   |   |   |   |   |   |   |   |
| CTTTGGCCAG GGTACGAAAG TTGAAATTAA ACGTACGGAA TTC  |   |   |   |   |   |   |   |   |   |   |   |   |   |
| GAAACCGGTC CCATGCTTTC AACTTTAATT TGCATGCCCTT AAG |   |   |   |   |   |   |   |   |   |   |   |   |   |
| BsiWI ECORI                                      |   |   |   |   |   |   |   |   |   |   |   |   |   |
| -----                                            |   |   |   |   |   |   |   |   |   |   |   |   |   |

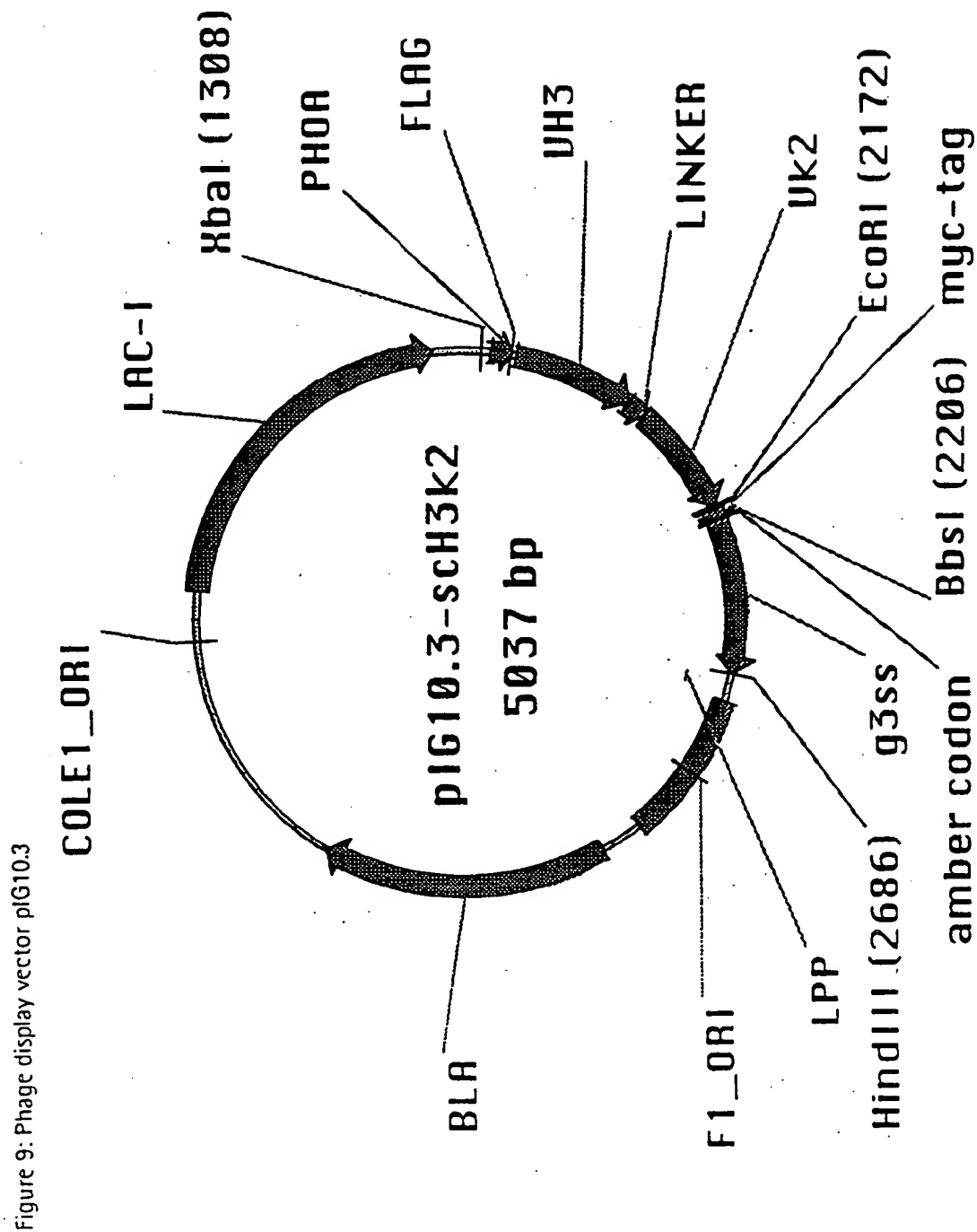


Figure 10: Sequence analysis of initial libraries

|   |      |   |   |   |   |   |   |   |   |   |   |   |   |
|---|------|---|---|---|---|---|---|---|---|---|---|---|---|
| A | 103  | W | W | W | W | W | W | W | W | W | W | W | W |
|   | 102  | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y |
|   | 101  | D | D | D | D | D | D | D | D | D | D | D | D |
|   | 100E | M | - | - | - | - | - | - | - | - | - | - | - |
|   | 100D | - | - | - | - | - | - | - | - | - | - | - | - |
|   | 100C | - | - | - | - | - | - | - | - | - | - | - | - |
|   | 100B | A | - | - | - | - | - | - | - | - | - | - | - |
|   | 100A | Y | - | - | - | - | - | - | - | - | - | - | - |
|   | 100  | F | Y | H | H | R | Y | P | - | S | K | A | D |
|   | 99   | G | N | W | Y | A | G | Q | R | N | S | A | Y |
|   | 98   | D | M | E | L | K | T | A | T | R | D | F | Q |
|   | 97   | G | K | T | E | L | T | E | I | N | G | T | P |
| B | 96   | G | G | R | R | F | N | N | A | Y | V | K | A |
|   | 95   | W | F | H | V | K | W | I | T | W | S | S | V |
|   | 94   | R | R | R | R | R | R | R | R | R | R | R | R |
|   | 93   | A | A | A | A | A | A | A | A | A | A | A | A |
|   | 92   | C | C | C | C | C | C | C | C | C | C | C | C |
|   |      |   |   |   |   |   |   |   |   |   |   |   |   |
|   |      |   |   |   |   |   |   |   |   |   |   |   |   |

Figure 10: Sequence analysis of initial libraries

|   |   |   |   |   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|---|---|---|---|
| C | C | C | C | C | C | C | C | C | C | C | C |
| A | A | A | A | A | A | A | A | A | A | A | A |
| R | R | R | R | R | R | R | R | R | R | R | R |
| Y | M | K | T | Y | * | R | M | K | S | Y |   |
| F | A | N | Q | P | G | N | K | G | W | A |   |
| V | L | Q | S | Y | S | P | P | S | T | G |   |
| H | R | M | F | R | G | W | M | E | N | T |   |
| F | A | V | W | S | S | N | L | F | D | T |   |
| L | S | F | E | N | E | V | N | L | K | F |   |
| Y | G | H | Q | F | H | N | R | E | P | K |   |
| T | K | A | Q | F | W | Y | D | T | N | Q |   |
| M | Y | R | K | M | S | L | G | D | F | G |   |
| V | I | K | V | P | I | H | T | V | I | P |   |
| M | M | F | M | M | F | F | M | M | M | M |   |
| D | D | D | D | D | D | D | D | D | D | D |   |
| V | V | V | Y | V | V | V | V | Y | V | Y |   |
| W | W | W | W | W | W | W | W | W | W | W |   |

Figure 11: Expression analysis of initial library

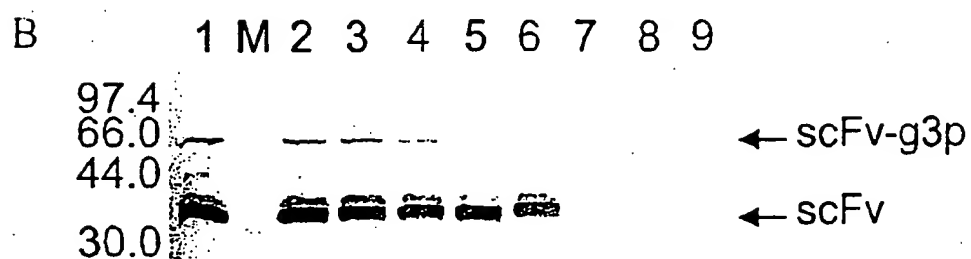
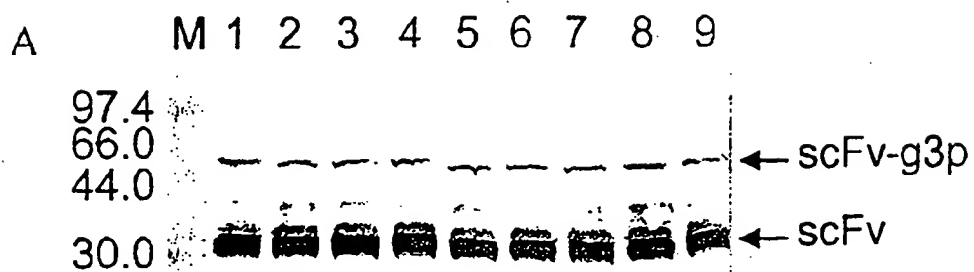




Figure 12: Increase of specificity during the panning rounds

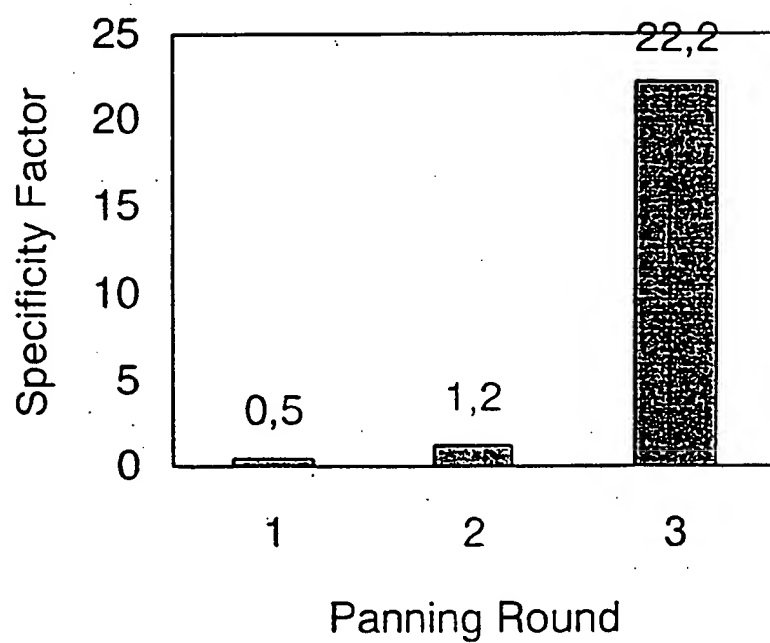


Figure 13: Phage ELISA of clones after the 3rd round of panning

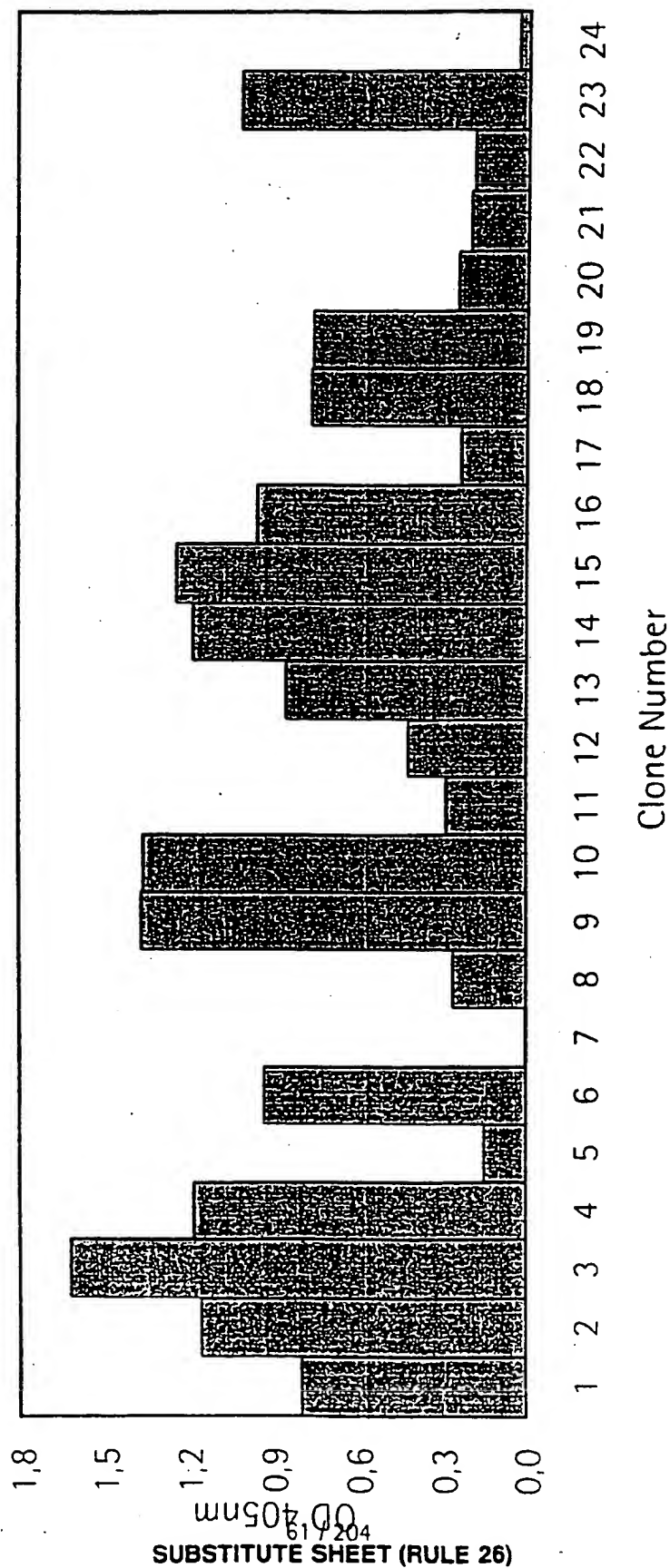


Figure 14: Competition ELISA

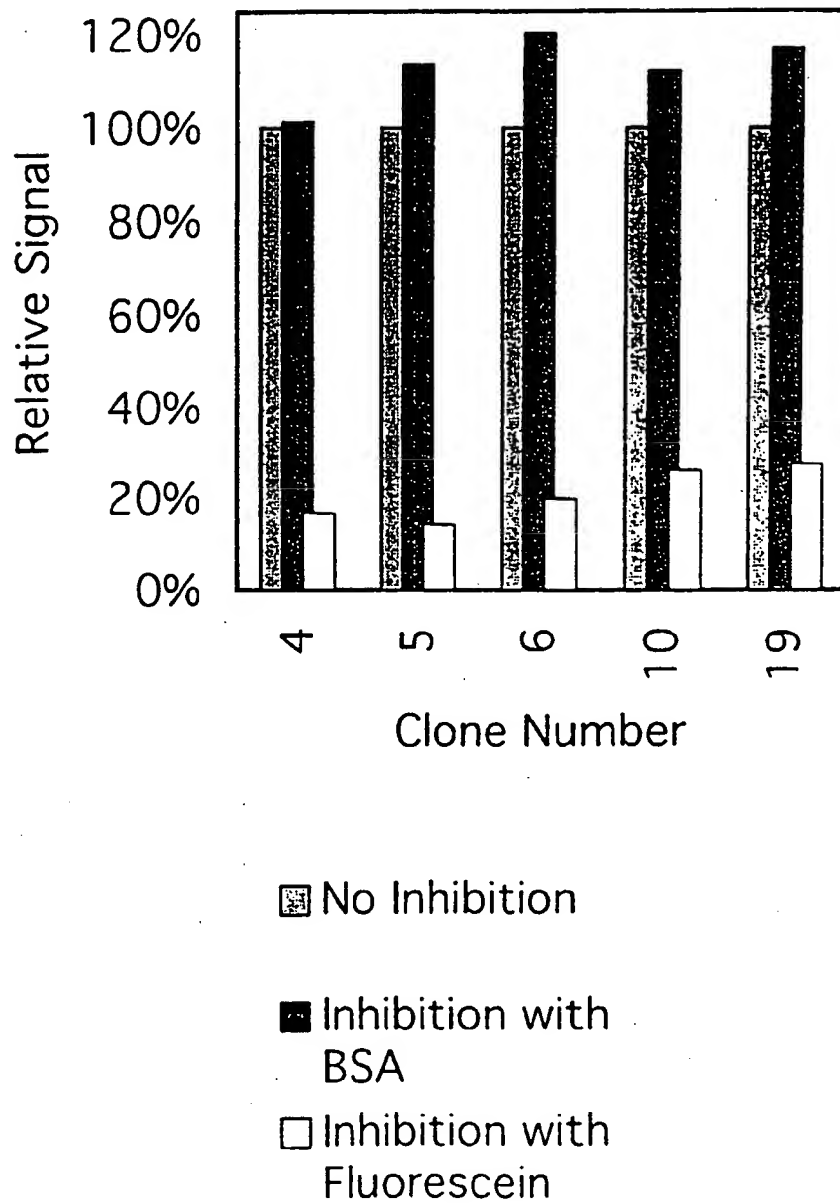


Figure 15: Sequence analysis of fluorescein binders

| Frequency | 1 | 3 | 1 | 2 | 1 | 1 | 1 | 2 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
|-----------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| 103       | W | W | W | W | W | W | W | W | W | W | W | W | W | W | W | W |
| 102       | > | > | > | > | > | > | > | Y | Y | > | > | > | > | > | > | Y |
| 101       | D | D | D | D | D | D | D | D | D | D | D | D | D | D | D | D |
| 100E      | F | F | F | F | F | M | F | F | F | F | F | F | F | F | F | F |
| 100D      | R | R | R | R | S | Q | V | K | Y | R | R | R | I | Q | R | R |
| 100C      | F | R | H | R | N | D | A | V | K | D | N | P | K | K | A | S |
| 100B      | R | M | R | K | K | F | K | T | V | M | M | R | R | R | F | F |
| 100A      | P | K | L | I | W | S | K | S | R | R | R | A | K | P | S | T |
| 100       | N | R | H | R | K | P | L | Y | S | R | G | F | G | Y | R | Y |
| 99        | Q | K | R | K | M | H | F | R | R | W | R | K | K | T | R | Q |
| 98        | N | Q | K | R | I | V | M | H | M | S | R | K | H | I | K | K |
| 97        | M | K | G | M | K | E | P | F | T | R | P | K | V | H | T | L |
| 96        | R | S | N | K | R | I | K | K | K | K | N | G | M | K | W | K |
| 95        | K | R | R | R | Y | L | R | R | R | K | R | K | R | R | R | K |
| 94        | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R |
| 93        | A | A | A | A | A | A | A | A | A | A | A | A | A | A | A | A |
| 92        | C | C | C | C | C | C | C | C | C | C | C | C | C | C | C | C |

Figure 16: Purification of fluorescein binding scFv fragments

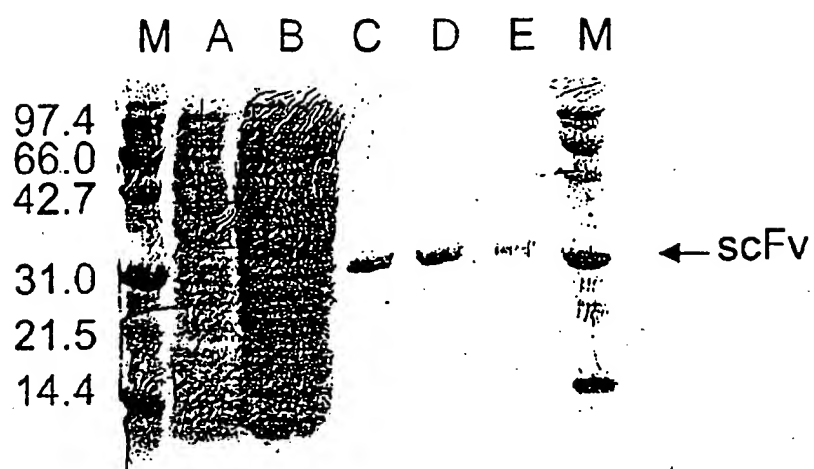


Figure 17: Enrichment factors after three rounds of panning

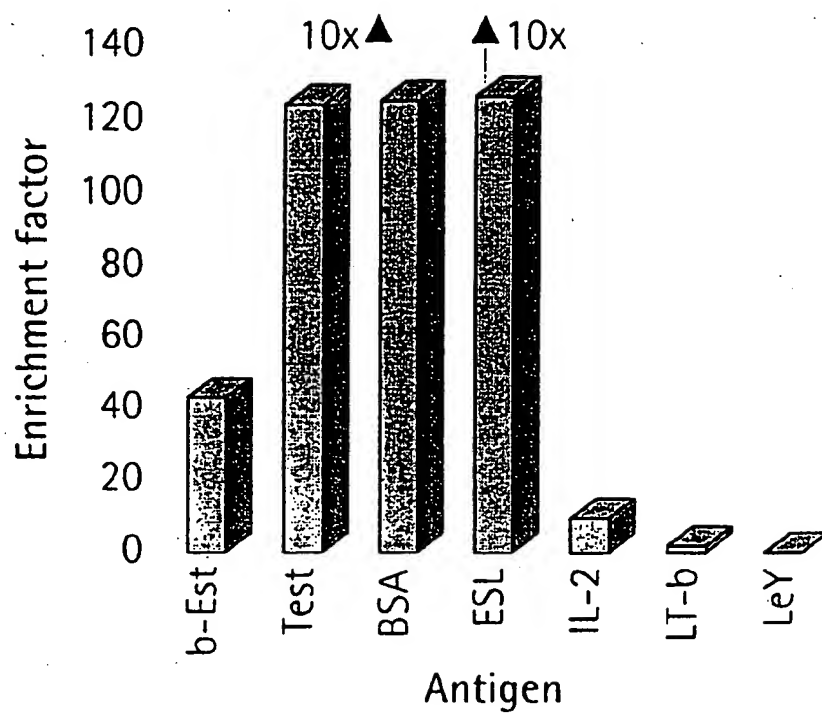


Figure 18: ELISA of anti-ESL-1 and anti- $\beta$ -estradiol antibodies

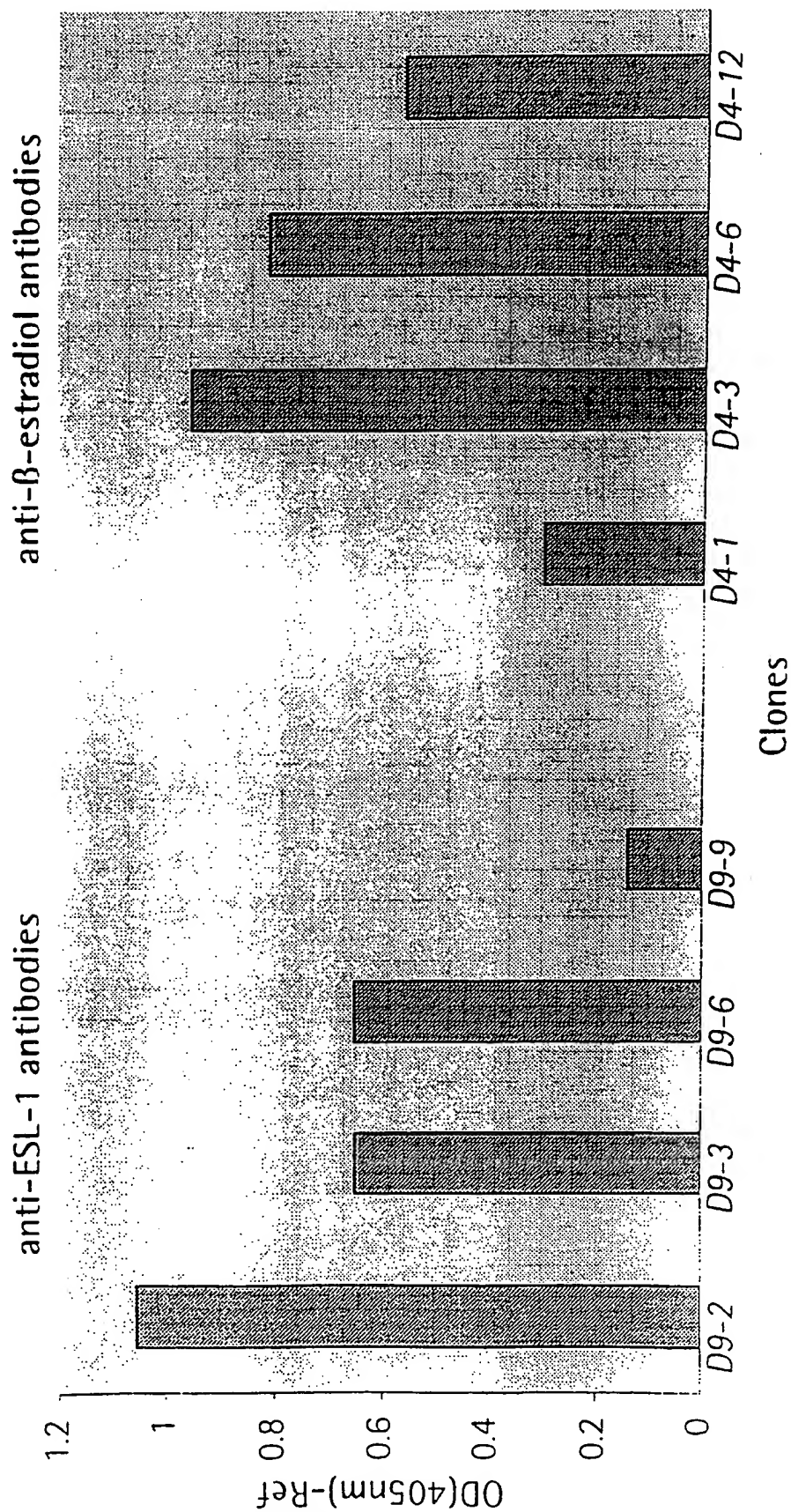


Figure 19: Selectivity and cross-reactivity of HuCAL antibodies

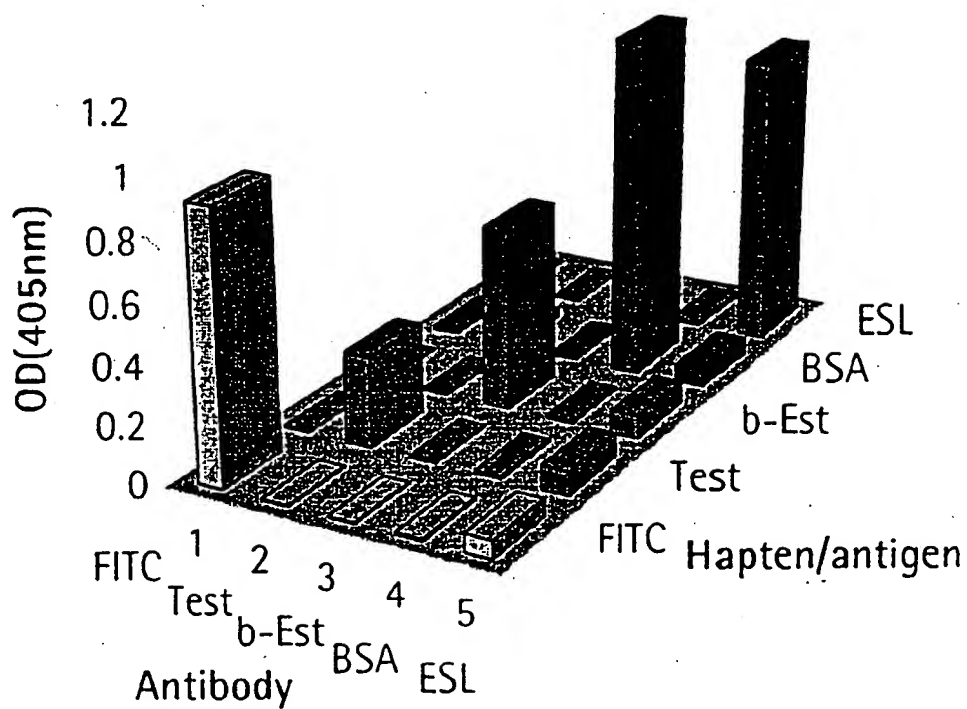




Figure 20: Sequence analysis of estradiol binders

| Frequency | 103 | 102 | 101 | 100E | 100D | 100C | 100B | 100A | 100 | 99 | 98 | 97 | 96 | 95 | 94 | 93 | 92 |
|-----------|-----|-----|-----|------|------|------|------|------|-----|----|----|----|----|----|----|----|----|
| 3         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 8         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 7         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 1         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 1         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 1         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 1         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 1         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 1         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 1         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 5         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 4         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |
| 1         | W   | W   | W   | W    | W    | W    | W    | W    | W   | W  | W  | W  | W  | W  | W  | W  | W  |

Figure 21: Sequence analysis of testosterone binders

|      | Frequency   |
|------|-------------|
| 103  | W W W W W W |
| 102  | Y Y Y V Y Y |
| 101  | D D D D D D |
| 100E | F F F F F F |
| 100D | A Q Q M W Q |
| 100C | L M M T K M |
| 100B | K K K K M Q |
| 100A | R Q N M I R |
| 100  | K W R W R S |
| 99   | A A A A R A |
| 98   | Q H Y G L R |
| 97   | K R K R P K |
| 96   | I N V K K R |
| 95   | Y Y Y Y R Y |
| 94   | R R R R R R |
| 93   | A A A A A A |
| 92   | C C C C C C |

Figure 22: Sequence analysis of lymphotoxin- $\beta$  binders

|      | Frequency |
|------|-----------|
| 103  | W 16      |
| 102  | V 1       |
| 101  | D 1       |
| 100E | F 1       |
| 100D | H 1       |
| 100C | G 1       |
| 100B | K 1       |
| 100A | I 1       |
| 100  | K 16      |
| 99   | S 1       |
| 98   | R 1       |
| 97   | Y 1       |
| 96   | R 1       |
| 95   | Q 1       |
| 94   | R 1       |
| 93   | A 1       |
| 92   | C 1       |

|   | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 | 24 | 25 | 26 | 27 | 28 | 29 | 30 | 31 | 32 | 33 | 34 | 35 | 36 | 37 | 38 | 39 | 40 | 41 | 42 | 43 | 44 | 45 | 46 | 47 | 48 | 49 | 50 | 51 | 52 | 53 | 54 | 55 | 56 | 57 | 58 | 59 | 60 | 61 | 62 | 63 | 64 | 65 | 66 | 67 | 68 | 69 | 70 | 71 | 72 | 73 | 74 | 75 | 76 | 77 | 78 | 79 | 80 | 81 | 82 | 83 | 84 | 85 | 86 | 87 | 88 | 89 | 90 | 91 | 92 | 93 | 94 | 95 | 96 | 97 | 98 | 99 | 100 |
|---|---|---|---|---|---|---|---|---|---|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|-----|
| 1 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 | 24 | 25 | 26 | 27 | 28 | 29 | 30 | 31 | 32 | 33 | 34 | 35 | 36 | 37 | 38 | 39 | 40 | 41 | 42 | 43 | 44 | 45 | 46 | 47 | 48 | 49 | 50 | 51 | 52 | 53 | 54 | 55 | 56 | 57 | 58 | 59 | 60 | 61 | 62 | 63 | 64 | 65 | 66 | 67 | 68 | 69 | 70 | 71 | 72 | 73 | 74 | 75 | 76 | 77 | 78 | 79 | 80 | 81 | 82 | 83 | 84 | 85 | 86 | 87 | 88 | 89 | 90 | 91 | 92 | 93 | 94 | 95 | 96 | 97 | 98 | 99 | 100 |

[illegible]

[illegible][illegible]

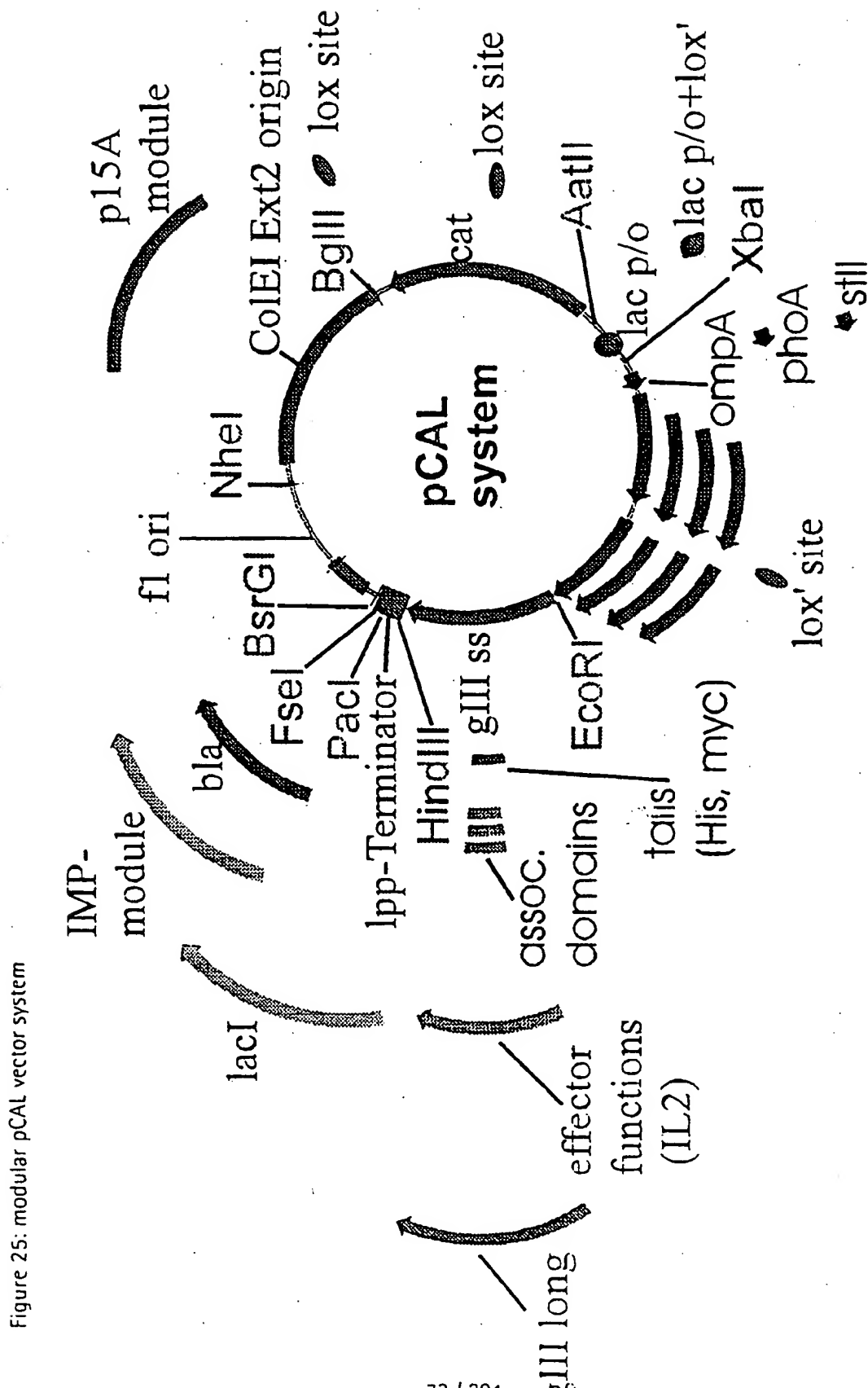


Figure 25: modular pCAL vector system

Figure 25a: List of unique restriction sites used in or suitable for HuCAL genes or pCAL vectors

| unique restriction site | Isoschizomers                     |
|-------------------------|-----------------------------------|
| AatII                   | /                                 |
| AfIII                   | BfrI, BspTI, Bst98I               |
| AscI                    | /                                 |
| Asel                    | Vspl, AsnI, PshBI                 |
| BamHI                   | BstI                              |
| BbeI                    | EheI, KasI, NarI                  |
| BbsI                    | BpuAI, BpiI                       |
| BglII                   | /                                 |
| BlpI                    | Bpu1102I, CelII, BliI             |
| BsaBI                   | MamI, Bsh1365I, BsrBRI            |
| BsiWI                   | Pfi23II, SphI, SnuI               |
| BspEI                   | AccIII, BseAI, BsiMI, Kpn2I, MroI |
| BsrGI                   | Bsp1407I, SspBI                   |
| BssHII                  | Paul                              |
| BstEII                  | BstPI, Eco91I, Eco065I            |
| BstXI                   | /                                 |
| Bsu36I                  | AocI, CvnI, Eco81I                |
| Drall                   | /                                 |
| DsmAI                   |                                   |
| EagI                    | BstZI, EclXI, Eco52I, XmaIII      |
| Eco57I                  | /                                 |
| EcoO109I                | Drall                             |
| EcoRI                   | /                                 |
| EcoRV                   | Eco32I                            |
| FseI                    | /                                 |
| HindIII                 | /                                 |
| HpaI                    | /                                 |
| KpnI                    | Acc65I, Asp718I                   |
| MluI                    | /                                 |
| MscI                    | BalI, MluNI                       |

Figure 25a: List of unique restriction sites used in or suitable for HuCAL genes or pCAL vectors

| unique restriction site | Isoschizomers                      |
|-------------------------|------------------------------------|
| MunI                    | MfeI                               |
| NheI                    | /                                  |
| NsiI                    | Ppu10I, EcoT22I, Mph1103I          |
| NspV                    | Bsp119I, BstBI, Csp45I, LspI, SfuI |
| PacI                    | /                                  |
| PmeI                    | /                                  |
| PmlI                    | BbrPI, Eco72I, PmaCI               |
| Psp5II                  | PpuMI                              |
| PstI                    | /                                  |
| RsrII                   | (RsrI), CpoI, CspI                 |
| SanDI                   | /                                  |
| SapI                    | /                                  |
| SexAI                   | /                                  |
| SpeI                    | /                                  |
| SfiI                    | /                                  |
| SphI                    | BbuI, PaeI, NspI                   |
| StuI                    | AatI, Eco147I                      |
| StyI                    | Eco130I, EcoT14I                   |
| XbaI                    | BspLU11II                          |
| XhoI                    | PaeR7I                             |
| XmaI                    | AvaI, SmaI, Cfr9I, PspAI           |



Figure 26: list of pCAL vector modules

| No   | module/flanking restriction sites | functional element                                            | sites to be removed | sites to be inserted | template      | reference                                                                                                                           |
|------|-----------------------------------|---------------------------------------------------------------|---------------------|----------------------|---------------|-------------------------------------------------------------------------------------------------------------------------------------|
| M1   | AatII-lacp/o-XbaI                 | lac promoter/operator                                         | 2x VspI (AseI)      | AatII                | vector pASK30 | Skerra et al. (1991) Bio/Technology 9, 273-278                                                                                      |
| M2   | BglII-lox-AatII                   | Cre/lox recombination site                                    | 2x VspI (AseI)      | lox, BglII           | (synthetic)   | Hoess et al. (1986) Nucleic Acids Res. 2287-2300                                                                                    |
| M3   | XbaI-lox'-SphI                    | Cre/lox' recombination site                                   | none                | lox', SphI           | (synthetic)   | see M2                                                                                                                              |
| M7-I | EcoRI-glllong-HindIII             | gllp of filamentous phage with N-terminal myctail/amber codon | SphI, BamHI         | none                 | vector pIG10  | Ge et al., (1994) Expressing antibodies in E. coli. In: Antibody engineering: A practical approach. IRL Press, New York, pp 229-266 |

Figure 26: list of pCAL vector modules

|         |                      |                                                                          |                            |                   |              |          |
|---------|----------------------|--------------------------------------------------------------------------|----------------------------|-------------------|--------------|----------|
| M7-II   | EcoRI-gIIIss-HindIII | truncated gIIIp of filamentous phage with N-terminal Gly-Ser linker      | SphI                       |                   | vector pIG10 | see M7-I |
| M7-III  | EcoRI-gIIIss-HindIII | truncated gIIIp of filamentous phage with N-terminal myctail/amber codon | SphI, BbsI                 |                   | vector pIG10 | see M7-I |
| M8      | SphI-lox-HindIII     | Cre/lox recombination site                                               | none                       | lox               | (synthetic)  | see M3   |
| M9-II   | HindIII-lpp-PacI     | lpp-terminator                                                           | none                       | PacI, FseI        | (synthetic)  | see M1   |
| M10-II  | PacI/FseI-bla-BsrGI  | beta-lactamase/bla (ampR)                                                | VspI, Eco57I, BssSI        | PacI, FseI, BsrGI | pASK30       | see M1   |
| M11-II  | BsrGI-f1 ori-NheI    | origin of single-stranded replication                                    | DrallI (BanII not removed) | BsrGI, NheI       | pASK30       | see M1   |
| M11-III | BsrGI-f1 ori-NheI    | origin of single-stranded replication                                    | DrallI, BanII              | BsrGI, NheI       | pASK30       | see M1   |

Figure 26: list of pCAL vector modules

|          |                      |                                                       |                            |                  |             |                                                                  |
|----------|----------------------|-------------------------------------------------------|----------------------------|------------------|-------------|------------------------------------------------------------------|
| M12      | NheI-p15A-BgIII      | origin of double-stranded replication                 | BssSI, VspI, NspV          | NheI, BgIII      | pACYC184    | Rose, R.E. (1988) Nucleic Acids Res. 16, 355                     |
| M13      | BgIII-lox-BgIII      | Cre/lox recombination site                            | none                       | BgIII, lox, XmnI | (synthetic) | see M3                                                           |
| M14-Ext2 | BgIII-ColEI-NheI     | origin of double-stranded replication                 | Eco57I (BssSI not removed) | BgIII, NheI      | pUC19       | Yanisch-Peron, C. (1985) Gene 33, 103-119                        |
| M17      | AatII-cat-BgIII      | chloramphenicol-acetyltransferase/cat (camR)          | BspEI, MscI, StyI/NcoI     |                  | pACYC184    | Cardoso, M. & Schwarz, S. (1992) J. Appl. Bacteriol. 72, 289-293 |
| M19      | XbaI-phoA-EcoRI      | signal sequence of phosphatase A                      | (synthetic)                |                  | (synthetic) | see M1                                                           |
| M20      | XbaI-phoA-FLAG-EcoRI | signal sequence of phosphatase A + FLAG detection tag | (synthetic)                |                  | (synthetic) | Knappik, A & Plückthun, A. (1994) BioTechniques 17, 754-761      |

Figure 26: list of pCAL vector modules

|     |                       |                                            |                                                    |  |             |                                                                               |
|-----|-----------------------|--------------------------------------------|----------------------------------------------------|--|-------------|-------------------------------------------------------------------------------|
| M21 | XbaI-stII-SapI        | heat-stable enterotoxin II signal sequence | (synthetic)                                        |  | (synthetic) | Lee et al. (1983) Infect. Immunol. 264-268                                    |
| M41 | AfIII-lacI-NheI       | lac-repressor                              | BstXI, MluI, BbsI, BanII, BstEII, HpaI, BbeI, VspI |  | pASK30      | see M1                                                                        |
| M42 | EcoRI-Histail-HindIII | poly-histidine tail                        | (synthetic)                                        |  | (synthetic) | Lindner et al., (1992) Methods: a companion to methods in enzymology 4, 41-56 |

Figure 27: functional map and sequence of MCS module

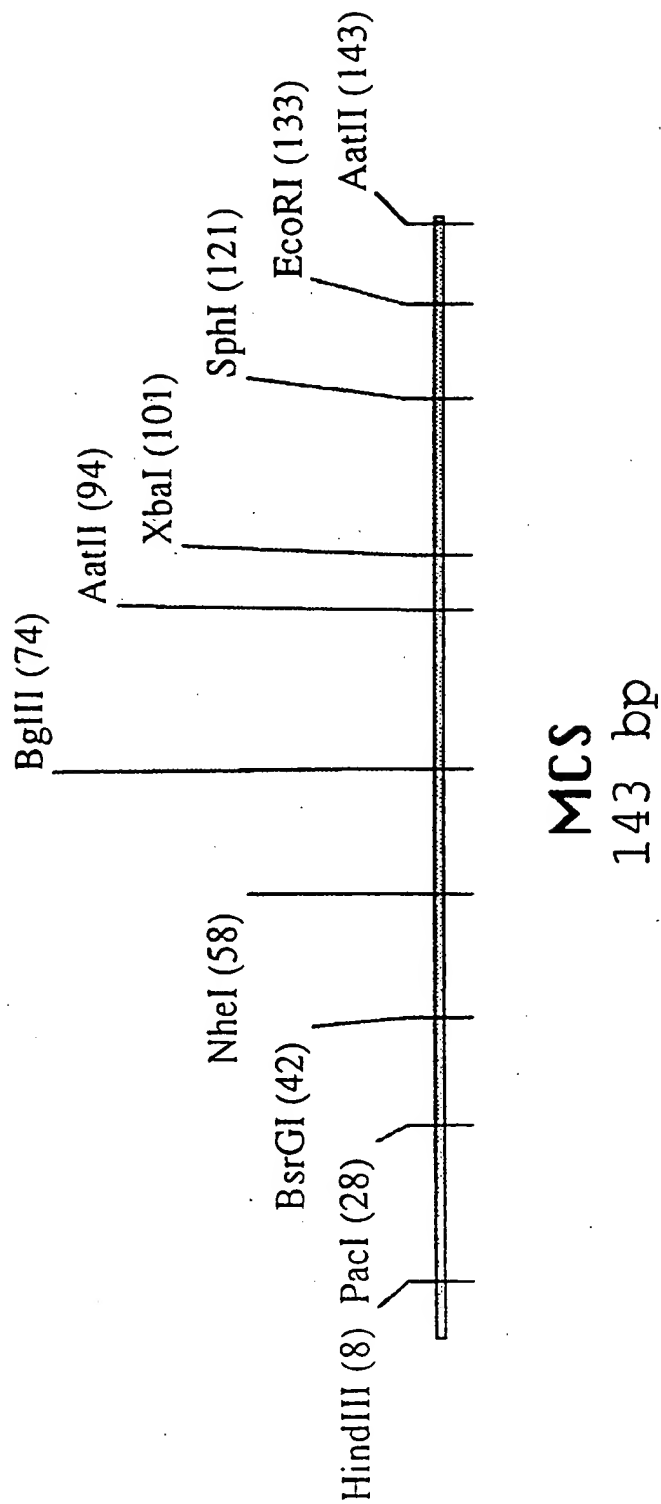


Figure 27: functional map and sequence of MCS module (continued)

|     | HindIII                                                 | PacI  | BsrGI       |
|-----|---------------------------------------------------------|-------|-------------|
|     | ~~~~~                                                   | ~~~~~ | ~~~~~       |
| 1   | ACATGTAAGC TTCCCCCCCC CCTTAATTAA CCCCCCCCCC TGTACACCCC  |       |             |
|     | TGTACATTCTG AAGGGGGGGG GGAATTAATT GGGGGGGGGG ACATGTGGGG |       |             |
|     |                                                         |       |             |
|     | NheI                                                    | BglII | AatII XbaI  |
|     | ~~~~~                                                   | ~~~~~ | ~~~~~       |
| 51  | CCCCCGCTA GCCCCCCCCC CCAGATCTCC CCCCCCCCCG GTCCCCCCT    |       |             |
|     | GGGGGGCGAT CGGGGGGGG GGTCTAGAGG GGGGGGGGCT GCAGGGGGGA   |       |             |
|     |                                                         |       |             |
|     | XbaI                                                    | SphI  | EcoRI AatII |
|     | ~~~~~                                                   | ~~~~~ | ~~~~~       |
| 101 | CTAGACCCCC CCCCCGCATG CCCCCCCCCC CGAATTCGAC GTC         |       |             |
|     | GATCTGGGGG GGGGGCGTAC GGGGGGGGGG GCTTAAGCTG CAG         |       |             |

Figure 28: functional map and sequence of pMCS cloning vector

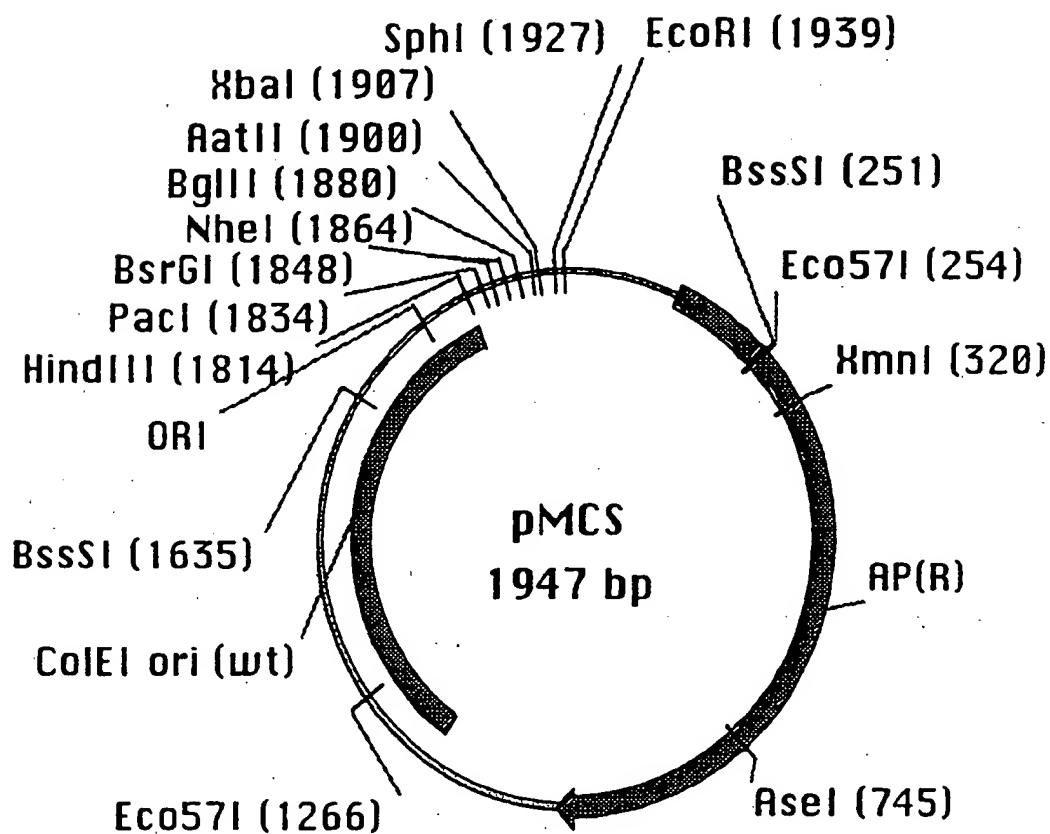


Figure 28: functional map and sequence of pMCS cloning vector (continued)

```

1  CAGGTGGCAC TTTTCGGGGA AATGTGCGCG GAACCCCTAT TTGTTTATTT
   GTCCACCGTG AAAAGCCCCCT TTACACGCGC CTTGGGGATA AACAAATAAA

51  TTCTAAATAC ATTCAAATAT GTATCCGCTC ATGAGACAAT AACCCTGATA
   AAGATTATG TAAGTTTATA CATAGGCGAG TACTCTGTTA TTGGGACTAT

101 AATGCTTCAA TAATATTGAA AAAGGAAGAG TATGAGTATT CAACATTGCC
   TTACGAAGTT ATTATAACTT TTTCCCTTCTC ATACTCATAA GTTGTAAGG

151 GTGTCGCCCT TATCCCTTT TTTGCGGCAT TTTGCCCTTC TGTTTTGTCT
   CACAGCGGGA ATAAGGGAAA AAACGCCGTA AAACGGAAGG ACAAAAACGA

                                     Eco57I
                                     ~~~~~

201 CACCCAGAAA CGCTGGTGAA AGTAAAAGAT GCTGAAGATC AGTTGGGTGC
 GTGGGTCTTT GCGACCACTT TCATTTTCTA CGACTTCTAG TCAACCCACG
 BSSI

251 ACGAGTGGGT TACATCGAAC TGGATCTCAA CAGCGGTAAG ATCCTTGAGA
 TGCTCACCCA ATGTAGCTTG ACCTAGAGTT GTCGCCATTC TAGGAACTCT
 BSSI
   ~~~~~

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Figure 28: functional map and sequence of pMCS cloning vector (continued)

XmnI

~~~~~

| | | | | | |
|-----|-------------|--------------|------------|------------|-------------|
| 301 | GTTTTCGCCCG | CGAAGAACGT | TTTCCAATGA | TGAGCACTTT | TAAAGTTCTG |
| | CAAAAGCGGG | GCTTCTTGCA | AAAGTTACT | ACTCGTGAAA | ATTTCAGAC |
| 351 | CTATGTGGCG | CGGTATTATC | CCGTATTGAC | GCCGGGCAAG | AGCAACTCGG |
| | GATACACCGC | GCCATAATAG | GGCATAACTG | CGCCCCGTTT | TCGTTGAGCC |
| 401 | TCGCCGCATA | CACTATTCTC | AGAAAGACTT | GGTTGAGTAC | TCACCAGTCA |
| | AGCGGCGTAT | GTGATAAGAG | TCTTACTGAA | CCAACTCATG | AGTGGTCAGT |
| 451 | CAGAAAAGCA | TCTTACGGAT | GGCATGACAG | TAAGAGAATT | ATGCAGTGCT |
| | GTCCTTTTCGT | AGAAATGCCCTA | CCGTACTGTC | ATTCTCTTAA | TACGTCACGA |
| 501 | GCCATAACCA | TGAGTGATAA | CACGCGGCC | AACTTACTTC | TGACAAACGAT |
| | CGGTATTGGT | ACTCACTATT | GTGACGCCCG | TTGAATGAAG | ACTGTTGCTA |
| 551 | CGGAGGACCG | AAGGAGCTAA | CCGCTTTTTT | GCACAACATG | GGGATCATG |
| | GCCTCCTGGC | TTCCCTCGATT | GGCGAAAAAA | CGTGTTGTAC | CCCCTAGTAC |
| 601 | TAACTCGCCT | TGATCGTTGG | GAACCGGAGC | TGAATGAAGC | CATACCAAAC |
| | ATTGAGCGGA | ACTAGCAACC | CTTGCCCTCG | ACTTACTTCG | GTATGGTTTG |
| 651 | GACGAGCGTG | ACACCACGAT | GCCTGTAGCA | ATGGCAACAA | CGTTGCGCAA |

Figure 28: functional map and sequence of pMCS cloning vector (continued)

| | | | | | |
|------|-------------|------------|-------------|-------------|-------------|
| | CTGCTCGCAC | TGTGGTGCTA | CGGACATCGT | TACCGTTGTT | GCAACGCGTT |
| | | | | AseI | |
| | | | | ~~~~~ | |
| 701 | ACTATTAACT | GGCGAACTAC | TTACTCTAGC | TTCCCCGGCAA | CAATTAATAG |
| | TGATAAATTGA | CCGCTTGATG | AATGAGATCG | AAGGGCCGTT | GTTAATTATC |
| 751 | ACTGGATGGA | GGCGGATAAA | GTTGCAGGAC | CAC TTCTGCG | CTCGGCCCTT |
| | TGACCTACCT | CCGCCATTAT | CAACGTCCCTG | GTGAAGACGC | GAGCCGGGAA |
| 801 | CCGGCTGGCT | GGTTTATTGC | TGATAAATCT | GGAGCCGGTG | AGCGTGGGTC |
| | GGCCGACCGA | CCAAATAACG | ACTATTTAGA | CCTCGGCCAC | TCGCACCCAG |
| 851 | TCGCGGTATC | ATTGCAGCAC | TGGGGCCAGA | TGGTAAGCCC | TCCC GTATCG |
| | AGCGCCATAG | TAACGTCGTG | ACCCCGGTCT | ACCATTGCGG | AGGCGATAGC |
| 901 | TAGTTATCTA | CACGACGGGG | AGTCAGGCAA | CTATGGATGA | ACGAAATAGA |
| | ATCAATAGAT | GTGCTGCCCC | TCAGTCCGTT | GATACCTACT | TGCTTTATCT |
| 951 | CAGATCGCTG | AGATAGGTGC | CTCACTGATT | AAGCATTTGGT | AACTGTCAGA |
| | GTCTAGCGAC | TCTATCCACG | GAGTGACTAA | TTCGTAACCA | TTGACAGTCT |
| 1001 | CCAAGTTTAC | TCATATATAC | TTTAGATTGA | TTTAAAACTT | CATTTTAAAT |
| | GGTTCAAATG | AGTATATATG | AAATCTAACT | AAATTTTGAA | GTAAAAATTA |

Figure 28: functional map and sequence of pMCS cloning vector (continued)

```

1051  TTAAAAGGAT CTAGGTGAAG ATCCTTTTGG ATAATCTCAT GACCAAAATC
      AATTTCCCTA GATCCACTTC TAGGAAAAC TATTAGAGTA CTGGTTTAG

1101  CCTTAACGTG AGTTTTCGTT CCACTGAGCG TCAGACCCCG TAGAAAAGAT
      GGAATGCGAC TCAAAAGCAA GGTGACTCGC AGTCTGGGC ATCTTTTCTA

1151  CAAAGGATCT TCTTGAGATC CTTTTTCTT GCGCGTAATC TGCTGCTTGC
      GTTCCCTAGA AGAACTCTAG GAAAAAAGA CCGCATTAG ACGACGAACG

1201  AAACAAAAAA ACCACCGCTA CCAGCGGTGG TTTGTTTGCC GGATCAAGAG
      TTTGTTT TGGTGCGGAT GGTCGCCACC AAACAAACGG CCTAGTTCTC

1251  CTACCAACTC TTTTCCGAA GGTAACGGC TTCAGCAGAG CGCAGATACC
      GATGGTTGAG AAAAAGGCTT CCATTGACCG AAGTCGTCTC GCGTCTATGG
                                     Eco57I
                                     ~~~~~~

1301  AAATACTGTC CTTCTAGTGT AGCCGTAGTT AGGCCACCAC TTCAAGAACT
      TTTATGACAG GAAGATCACA TCGGCATCAA TCCGGTGGTG AAGTTCTTGA

1351  CTGTAGCACC GCCTACATAC CTCGCTCTGC TAATCCTGTT ACCAGTGGCT
      GACATCGTGG CGGATGTATG GAGCGAGACG ATTAGGACAA TGGTCACCGA

```

Figure 28: functional map and sequence of pMCS cloning vector (continued)

```

1401  GCTGCCAGTG  GCGATAAGTC  GTGTCTTACC  GGGTTGGACT  CAAGACGATA
      CGACGGTCAC  CGCTATTTCAG  CACAGAATGG  CCCAACCTGA  GTTCTGCTAT

1451  GTTACCGGAT  AAGGCGCAGC  GGTCGGGCTG  AACGGGGGGT  TCGTGCCACAC
      CAATGGCCCTA  TTCCGCGGTCG  CCAGCCCCGAC  TTGCCCCCCCA  AGCACGTGTG

1501  AGCCCCAGCTT  GGAGCGAACG  ACCTACACCG  AACTGAGATA  CCTACAGCGT
      TCGGGTCGAA  CCTCGCTTGC  TGGATGTGGC  TTGACTCTAT  GGATGTGCGA

1551  GAGCTATGAG  AAAGCGC  AC  GCTTCCCGAA  GGGAGAAAGG  CGGACAGGTA
      CTCGATACTC  TTTCGCGGTG  CGAAGGGCTT  CCTCTTTCC  GCCTGTCCAT

1601  TCCGGTAAGC  GGCAGGGTCG  GAACAGGAGA  GCGCACGAGG  GAGCTTCCAG
      AGGCCATTTCG  CCGTCCCAGC  CTTGTCCCTCT  CGCGTGCTCC  CTCGAAAGTC

      BssSI
      ~~~~~

1651  GGGGAAACGC  CTGGTATCTT  TATAGTCCTG  TCGGGTTTCC  CCACCTCTGA
      CCCCCTTTGCG  GACCATAGAA  ATATCAGGAC  AGCCCCAAAGC  GGTGGAGACT

1701  CTTGAGCGTC  GATTTTGTG  ATGCTCGTCA  GGGGGGCGGA  GCCTATGGAA
      GAACTCGCAG  CTAAAAACAC  TACGAGCAGT  CCCCCCGCCT  CGGATACCTT

1751  AAACGCCAGC  AACGCGGCCT  TTTTACGGTT  CCTGGCCCTT  TGCTGGCCCTT

```

Figure 28: functional map and sequence of pMCS cloning vector (continued)

| | | | | | |
|------|------------|------------|-------------|-------------|------------|
| | TTTGCGGTCG | TTGCGCCGGA | AAATGCCAA | GGACCGGAA | ACGACCGGAA |
| | | HindIII | | PacI | BsrGI |
| | | ~~~~~ | | ~~~~~ | ~~~~~ |
| 1801 | TTGCTCACAT | GTAAGCTTCC | CCCCCCCCTT | AATTAACCC | CCCCCCTGTA |
| | AACGAGTGTA | CATTCGAAGG | GGGGGGGAA | TTAATTGGG | GGGGGACAT |
| | | NheI | | BglII | AatII |
| | | ~~~~~ | | ~~~~~ | ~~~~~ |
| 1851 | CACCCCCCCC | CCGCTAGCCC | CCCCCCCCCAG | ATCTCCCCC | CCCCGACGTC |
| | GTGGGGGGGG | GGCGATCGGG | GGGGGGGGTC | TAGAGGGGG | GGGGCTGCAG |
| | | XbaI | | SphI | EcoRI |
| | | ~~~~~ | | ~~~~~ | ~~~~~ |
| 1901 | CCCCCTCTAG | ACCCCCCCCC | CGCATGCCCC | CCCCCCCCGAA | TTCACGT |
| | GGGGGAGATC | TGGGGGGGGG | GCGTACGGGG | GGGGGGGCTT | AAGTGCA |

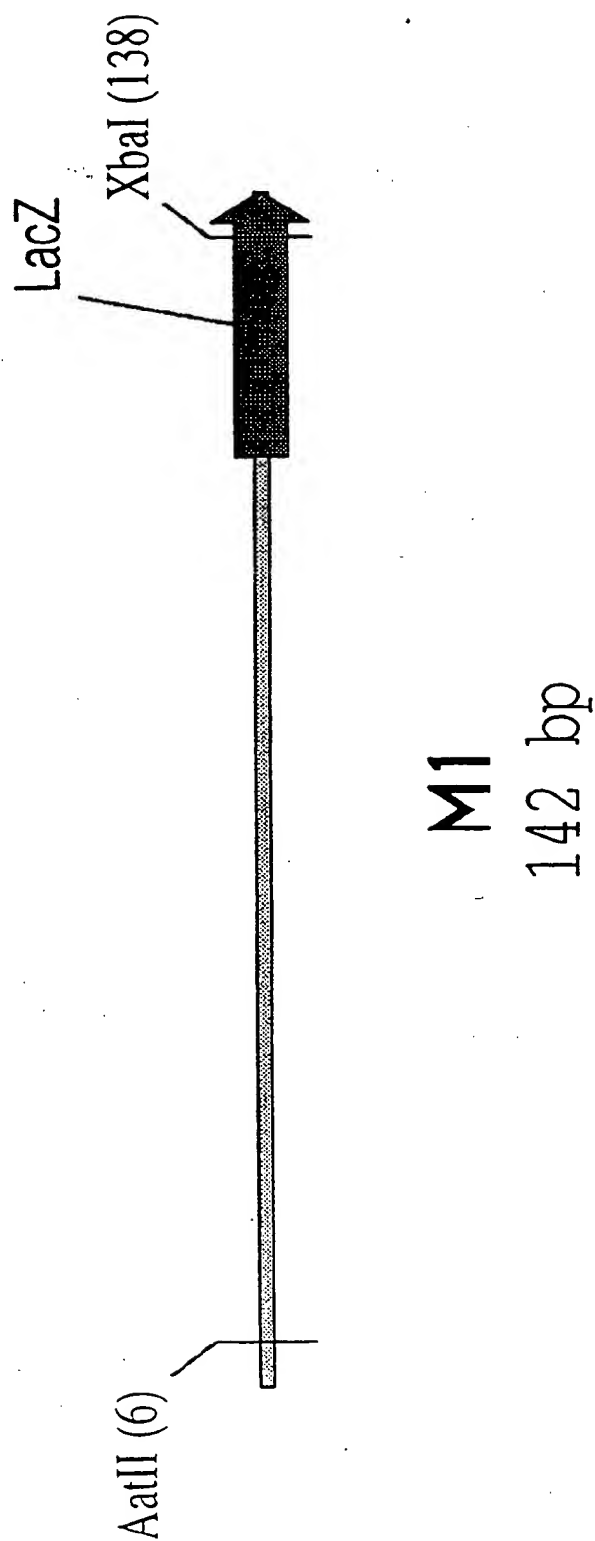


Figure 29: functional map and sequence of pCAL module M1

Figure 29: functional map and sequence of pCAL module M1

AatII
~~~~~  
1 GACGTCTTAA TGTGAGTTAG CTCACTCATT AGGCACCCCA GGCTTTACAC  
CTGCAGAATT ACACTCAATC GAGTGAGTAA TCCGTGGGGT CCGAAATGTG  
  
51 TTTATGCTTC CGGCTCGTAT GTTGTGTGGA ATTGTGAGCG GATAACAATT  
AAATACGAAG GCCGAGCATA CAACACACCT TAACACTCGC CTATTGTTAA  
  
XbaI  
~~~~~  
101 TCACACAGGA AACAGCTATG ACCATGATTA CGAATTCTA GA
AGTGTGTCCT TTGTCGATAC TGGTACTAAT GCTAAAGAT CT

Figure 30: functional map and sequence of pCAL module M7-II

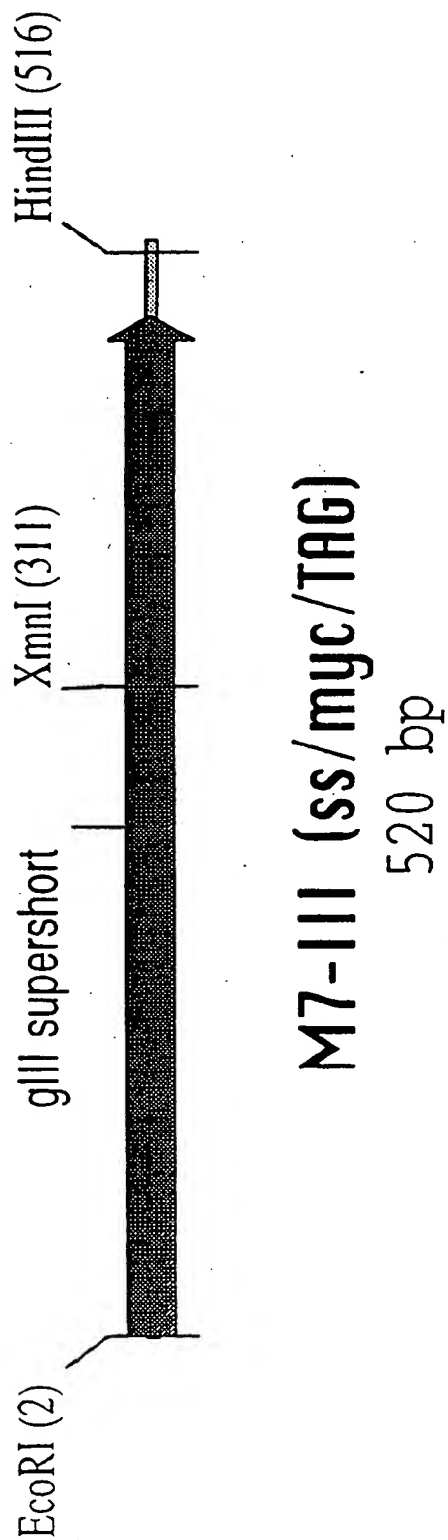


Figure 30: functional map and sequence of pCAL module M7-II (continued)

EcoRI
 ~~~~~  
 1 GAATTCGAGC AGAAGCTGAT CTCTGAGGAG GATCTGTAGG GTGGTGGCTC  
 CTTAAGCTCG TCTTCGACTA GAGACTCCTC CTAGACATCC CACCACCGAG

51 TGGTTCGGGT GATTTTGATT ATGAAAAGAT GGCAAACGCT AATAAGGGGG  
 ACCAAGGCCA CTAAAACTAA TACTTTTCTA CCGTTTGCGA TTATTCCCCC

101 CTATGACCGA AAATGCCGAT GAAAACGCCG TACAGTCTGA CGCTAAAGGC  
 GATACTGGCT TTTACGGGCTA CTTTGGCGCG ATGTCAGACT GCGATTTCGG

151 AAACCTTGATT CTGTCGGCTAC TGATTACGGT GCTGCTATCG ATGGTTTCAT  
 TTTGAACTAA GACAGCGATG ACTAATGCCA CGACGATAGC TACCAAAGTA

201 TGGTGACGTT TCCGGCCTTG CTAATGGTAA TGGTGCTACT GGTGATTTTG  
 ACCACTGCAA AGGCCGGAAC GATTACCATT ACCACGATGA CCACTAAAC

251 CTGGCTCTAA TTCCCAAATG GCTCAAGTCG GTGACGGTGA TAATTCACCT  
 GACCGAGATT AAGGGTTTAC CGAGTTCAGC CACTGCCACT ATTAAGTGGA

XmnI  
 ~~~~~  
 301 TTAATGAATA ATTTCCGTCA ATATTACCT TCCCTCCCTC AATCGGTGA
 AATTACTTAT TAAAGGCAGT TATAAATGGA AGGAGGGAG TTAGCCAACT

Figure 30: functional map and sequence of pCAL module M7-II (continued)

```
351  ATGTCGCCCT  TTTGTCTTTG  GCGCTGGTAA  ACCATATGAA  TTTTCTATTG
      TACAGCGGGA  AACAGAAAC  CGCGACCATT  TGGTATACTT  AAAAGATAAC

401  ATTGTGACAA  AATAAACTTA  TTCCGTGGTG  TCTTTGCCGT  TCTTTTATAT
      TAACACTGTT  TTATTGAAT  AAGGCACCAC  AGAAACGCAA  AGAAATATA

451  GTTGCCACCT  TTATGTATGT  ATTTCTACG  TTGCTAACA  TACTGCGTAA
      CAACGGTGA  AATACATACA  TAAAGATGC  AAACGATTGT  ATGACGCATT

501  TAAGGAGTCT  TGATAAGCTT
      ATTCCTCAGA  ACTATTCCGAA
```

HindIII

~~~~~

Figure 31: functional map and sequence of pCAL module M9-II

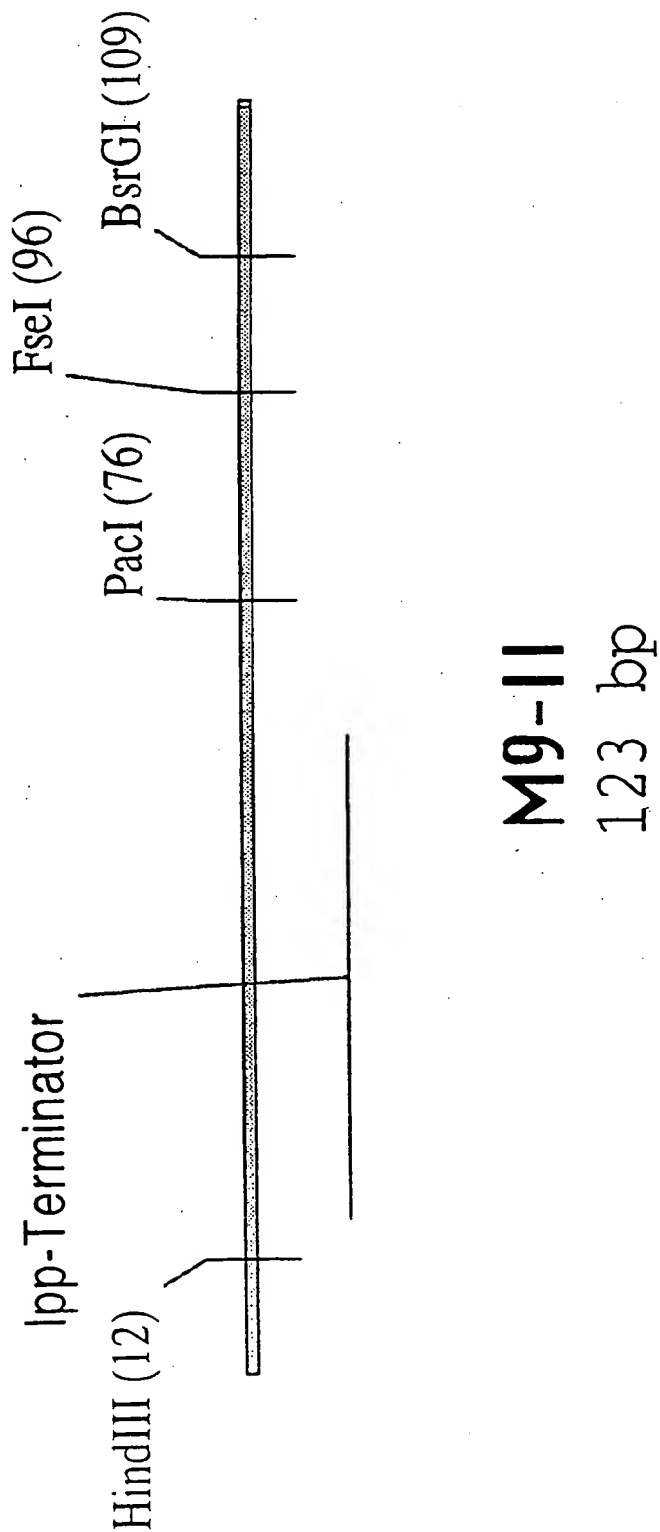


Figure 31: functional map and sequence of pCAL module M9-II (continued)

```

HindIII
~~~~~
1 GGGGGGGGGG AAGCTTGACC TGTGAAGTGA AAAATGGCGC AGATTGTGCG
 CCCCCCCCCC TTCGAACTGG ACACTTCACT TTTTACCGCG TCTAACACGC

PacI
~~~~~
51 ACATTTT TTTT TGCTGCCGT TTAATTAAAG GGGGGGGGGG GCCGGCCTGG
   TGTAATAAAA ACAGACGGCA AATTAATTTC CCCCCCCCCC CGCCCGGACC

BsrGI
~~~~~
101 GGGGGGGTGT ACAGGGGGG GGG
 CCCCCCCACA TGTCCCCCCC CCC

```

Figure 32: functional map and sequence of pCAL module M11-III

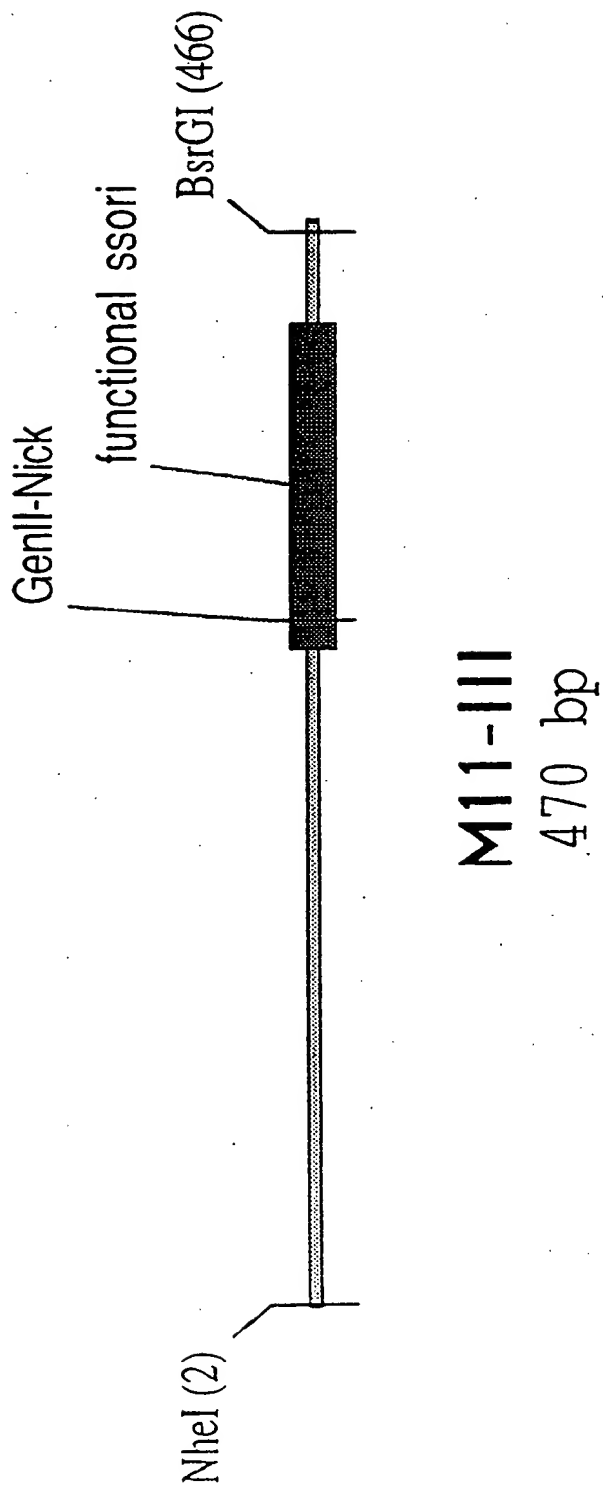


Figure 32: functional map and sequence of pCAL module M11-III (continued)

NheI

~~~~~

1	GCTAGCACGC	GCCCTGTAGC	GGCGCATTA	GGCGGCGGG	TGTGGTGGTT
	CGATCGTGCG	CGGACATCG	CCGCGTAATT	CGCGCCGCC	ACACCACCAA
51	ACGCGCAGCG	TGACCGCTAC	ACTTGCCAGC	GCCCTAGCGC	CCGCTCCTTT
	TGCGCGTCCG	ACTGGCGATG	TGAACGGTCG	CGGATCGCG	GGCGAGGAAA
101	CGCTTTCTTC	CCTTCCTTTC	TCGCCACGTT	CGCCGGCTTT	CCCCGTCAAG
	GCGAAAGAAG	GGAAGGAAAG	AGCGGTGCAA	GCGGCCGAAA	GGGCGAGTTC
151	CTCTAAATCG	GGGCATCCCT	TTAGGGTTCC	GATTTAGTGC	TTTACGGCAC
	GAGATTTAGC	CCCGTAGGGA	AATCCCAAGG	CTAAATCACG	AAATGCCGTG
201	CTCGACCCCA	AAAACTTGA	TTAGGGTGAT	GGTTCTCGTA	GTGGGCCATC
	GAGCTGGGGT	TTTTTGAAC	AATCCCACTA	CCAAGAGCAT	CACCCGGTAG
251	GCCCTGATAG	ACGGTTTTC	GCCCTTTGAC	GTGGAGTCC	ACGTTCTTTA
	CGGACTATC	TGCCAAAAG	CGGAAACTG	CAACCTCAGG	TGCAAGAAAT
301	ATAGTGGA	CTTGTTCCAA	ACTGGAACAA	CACCTAACCC	TATCTCGGTC
	TATCACCTGA	GAACAAGGT	TGACCTTGTT	GTGAGTTGGG	ATAGAGCCAG
351	TATTCCTTTG	ATTATAAGG	GATTTGCCG	ATTCGGCCT	ATTGGTTAAA

Figure 32: functional map and sequence of pCAL module M11-III (continued)

	ATAAGAAAAC	TAAATATTCC	CTAAAACGGC	TAAAGCCGGA	TAAACCAATTT
401	AAATGAGCTG	ATTTAACAAA	AATTTAACGC	GAATTTTAAC	AAAATATTAA
	TTTACTCGAC	TAAATTGTTT	TTAAATTGCG	CTTAAAATTG	TTTATATAATT
			BsrgI		
			~~~~~		
451	CGTTTACAAT	TTCATGTACA			
	GCAAATGTTA	AAGTACATGT			

Figure 33: functional map and sequence of pCAL module M14-Ext2

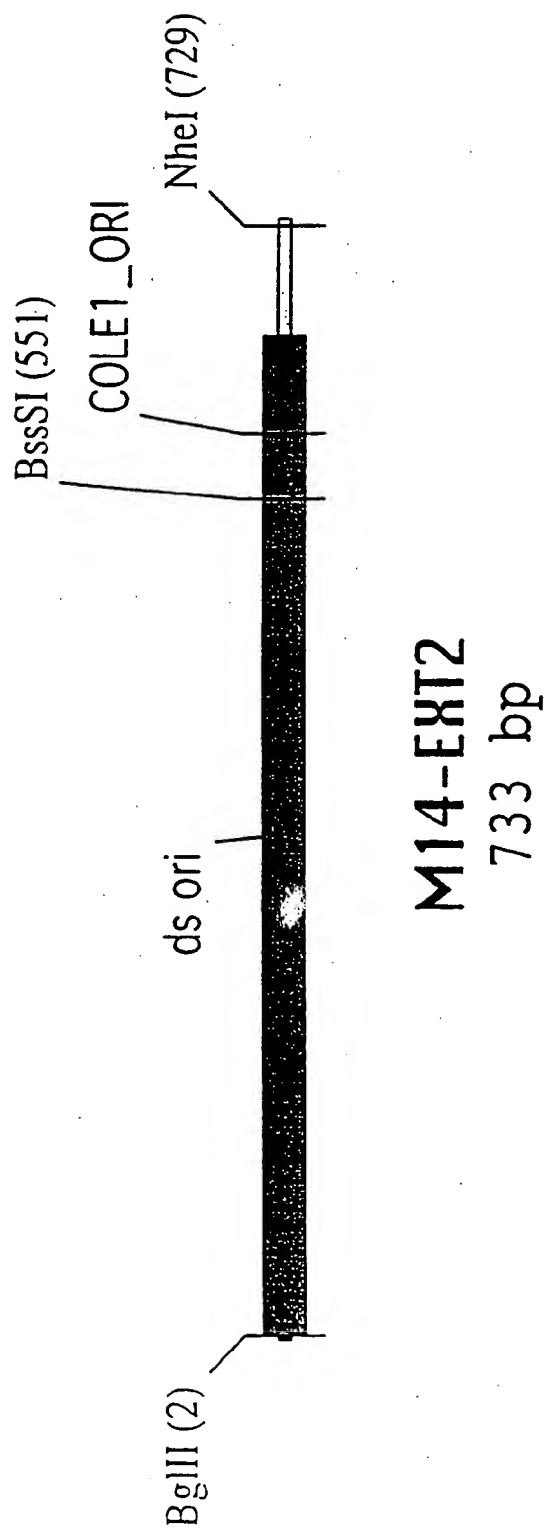




Figure 33: functional map and sequence of pCAL module M14-Ext2 (continued)

BglII  
~~~~~

|     |            |            |            |             |             |
|-----|------------|------------|------------|-------------|-------------|
| 1   | AGATCTGACC | AAAATCCCTT | AACGTGAGTT | TTCGTTCCAC  | TGAGCGTCAG  |
|     | CTAGACTGG  | TTTTAGGGAA | TTGCACTCAA | AAGCAAGGTG  | ACTCGCAGTC  |
| 51  | ACCCCGTAGA | AAAGATCAAA | GGATCTTCTT | GAGATCCCTT  | TTTTCTGCGC  |
|     | TGGGGCATCT | TTTCTAGTTT | CCTAGAAGAA | CTCTAGGAAA  | AAAAGACGCG  |
| 101 | GTAATCTGCT | GCTTGCAAAC | AAAAAACA   | CCGCTACCAG  | CGGTGGTTTG  |
|     | CATTAGACGA | CGAACGTTTG | TTTTTTTGGT | GGCGATGGTC  | GCCACCAAAC  |
| 151 | TTTGCCGGAT | CAAGAGCTAC | CAACTCTTTT | TCCGAAGGTA  | ACTGGCTACA  |
|     | AAACGGCCTA | GTTCTCGATG | GTTGAGAAAA | AGGCTTCCAT  | TGACCGATGT  |
| 201 | GCAGAGCGCA | GATACCAAAT | ACTGTTCTTC | TAGTGTAGCC  | GTAGTTAGGC  |
|     | CGTCTCGCGT | CTATGGTTTA | TGACAAGAAG | ATCACATCCG  | CATCAATCCG  |
| 251 | CACCACTTCA | AGAACTCTGT | AGCACCGCCT | ACATACCCTCG | CTCTGCTAAT  |
|     | GTGGTGAAGT | TCTTGAGACA | TCGTGGCGGA | TGTATGGAGC  | GAGACGATTA  |
| 301 | CCTGTTACCA | GTGGCTGCTG | CCAGTGCGCA | TAAGTCGTGT  | CTTACCGGGT  |
|     | GGACAATGGT | CACCGACGAC | GGTCACCGCT | ATTCAGCACA  | GAATGGCCCCA |
| 351 | TGGACTCAAG | ACGATAGTTA | CCGGATAAGG | CGCAGCGGTC  | GGGCTGAACG  |

Figure 33: functional map and sequence of pCAL module M14-Ext2 (continued)

```

ACCTGAGTTC TGCTATCAAT GGCCATTCC GCGTCGCCAG CCCGACTTGC
401 GGGGGTTCGT GCACACAGCC CAGCTTGGAG CGAACGACCT ACACCGAACT
CCCCAAGCA CGTGTGTCGG GTCGAACCTC GCTTGCTGGA TGTGGCTTGA

451 GAGATACCTA CAGCGTGAGC TATGAGAAAG CGCCACGCTT CCCGAAGGGA
CTCTATGGAT GTCGCACTCG ATACTCTTTC GCGGTGCGAA GGGCTTCCCT

501 GAAAGGCGGA CAGGTATCCG GTAAGCGGCA GGTTCGGAAC AGGAGAGCGC
CTTCCGCCT GTCCATAGGC CATTCGCCGT CCCAGCCTTG TCCTCTCGCG
BssSI

551 ACGAGGGAGC TTCCAGGGGG AAACGCCCTGG TATCTTTATA GTCCGTGTCGG
TGCTCCCTCG AAGTCCCCC TTGCGGACC ATAGAAATAT CAGGACAGCC
BssSI
~~~~~

601 GTTTCGCCAC CTCGACTTG AGCGTCGATT TTTGTGATGC TCGTCAGGGG
CAAAGCGGTG GAGACTGAAC TCGCAGCTAA AAACACTACG AGCAGTCCCC

651 GGCGGAGCCT ATGGAAAAAC GCCAGCAACG CGGCCCTTTT ACGGTTCCCTG
CCGCCCTCGGA TACCTTTTTC CGGTCGTTCG GCCGGAAAAA TGCCAAGGAC

```

Figure 33: functional map and sequence of pCAL module M14-Ext2 (continued)

NheI  
~~~~~  
701 GCCTTTTGCT GGCCTTTTGC TCACATGGCT AGC
CGGAAAACGA CCGGAAAACG AGTGTAACCGA TCG

Figure 34: functional map and sequence of pCAL module M17

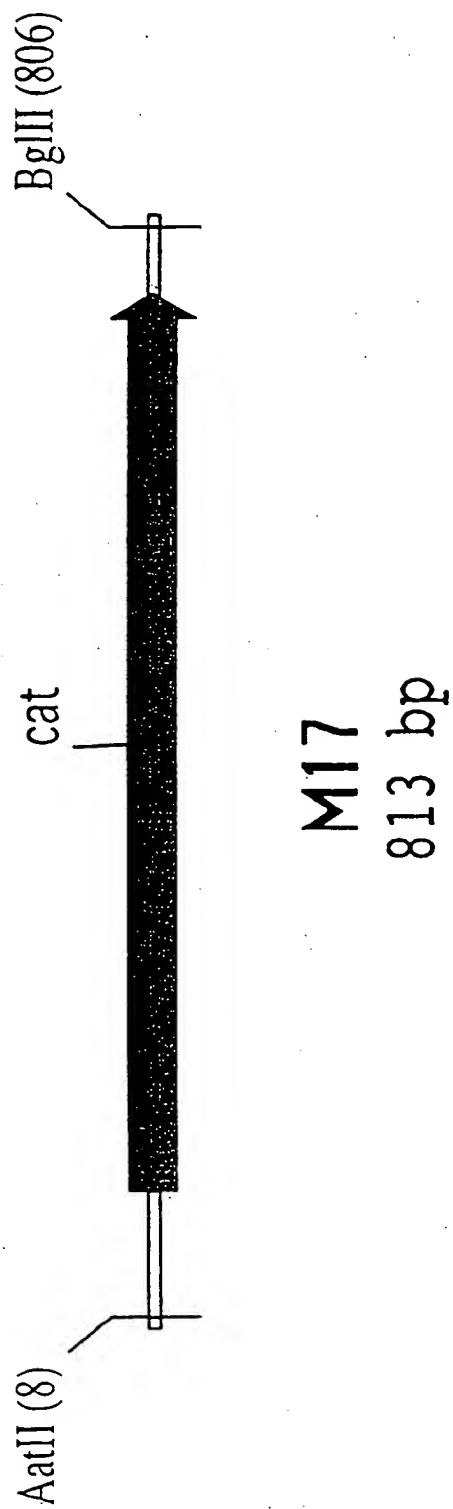


Figure 34: functional map and sequence of pCAL module M17 (continued)

AatII

~~~~~

|     |            |              |             |             |              |
|-----|------------|--------------|-------------|-------------|--------------|
| 1   | GGGACGTCGG | GTGAGGTTCC   | AAC TTCACC  | ATAATGAAAT  | AAGATCACTA   |
|     | CCCTGCAGCC | CACTCCAAGG   | TTGAAAGTGG  | TATTA CTTTA | TTCTAGTGAT   |
| 51  | CCGGGCGTAT | TTTTTGAGTT   | ATCGAGATT   | TCAGGAGCTA  | AGGAAGCTAA   |
|     | GGCCCCGATA | AAA AACTCAA  | TAGCTCTAAA  | AGTCC TCGAT | TCCTTCGATT   |
| 101 | AATGGAGAAA | AAAATCACTG   | GATATACCAC  | CGTTGATATA  | TCCCAATGGC   |
|     | TTACCTCTTT | TTTTAGTGAC   | CTATATGGTG  | GCAACTATAT  | AGGTTACCG    |
| 151 | ATCGTAAAGA | ACATTTTGAG   | GCAT TTCAGT | CAGTTGCTCA  | ATGTACCTAT   |
|     | TAGCATTTCT | TGTA AAACTC  | CGTAAAGTCA  | GTC AACGAGT | TACATGGATA   |
| 201 | AACCAGACCG | TTCAGCTGGA   | TATTACGGCC  | TTTTTAAAGA  | CCGTAAAGAA   |
|     | TTGGTCTGGC | AAGTCGACCT   | ATAATGCCCG  | AAAAATTTCT  | GGCATTTCTT   |
| 251 | AAATAAGCAC | AAGTTT TATC  | CGGCCTTTAT  | TCACAT TCTT | GCCCGCCCTGA  |
|     | TTTATTCGTG | TTCA AAAATAG | GCCGGAAATA  | AGTGTAAGAA  | CGGGCGGACT   |
| 301 | TGAATGCTCA | CCCGGAGTTC   | CGTATGGCAA  | TGAAAGACGG  | TGAGCTGGTG   |
|     | ACTTACGAGT | GGGCTCAAG    | GCATACCGTT  | ACTTTCTGCC  | ACTCGACCAC   |
| 351 | ATATGGGATA | GTGTTACCC    | TTGTTACACC  | GTTTTC CATG | AGCAA AACTGA |

Figure 34: functional map and sequence of pCAL module M17 (continued)

|     |            |            |            |             |             |
|-----|------------|------------|------------|-------------|-------------|
|     | TATACCCTAT | CACAAGTGGG | AACAATGTGG | CAAAAGGTAC  | TCGTTTGA    |
| 401 | AACGTTTTC  | TCGCTCTGA  | GTGAATACCA | CGACGATTC   | CGGCAGTTTC  |
|     | TTGCAAAAGT | AGCGAGACCT | CACTTATGGT | GCTGCTAAAG  | GCCGTCAAAG  |
| 451 | TACACATATA | TTCGCAAGAT | GTGGCGTGT  | ACGGTGAAA   | CCTGGCCTAT  |
|     | ATGTGTATAT | AAGCGTTCTA | CACCGCACAA | TGCCACTTTT  | GGACCCGATA  |
| 501 | TTCCCTAAAG | GGTTTATTGA | GAATATGTTT | TTCGTCTCAG  | CCAATCCCCTG |
|     | AAGGGATTTC | CCAAATAACT | CTTATACAAA | AAGCAGAGTC  | GGTAGGGAC   |
| 551 | GGTGAGTTTC | ACCAGTTTGT | ATTTAAACGT | AGCCAATATG  | GACAACTTCT  |
|     | CCACTCAAAG | TGGTCAAAAC | TAAATTGCA  | TCCGTTATAC  | CTGTTGAAGA  |
| 601 | TCGCCCCCGT | TTTCACTATG | GGCAAATAT  | ATACGCAAGG  | CGACAAGGTG  |
|     | AGCGGGGGCA | AAAGTGATAC | CCGTTTATAA | TATGCGTTCC  | GCTGTTCCAC  |
| 651 | CTGATGCCGC | TGGCGATTCA | GGTTCATCAT | GCCGTTTGTG  | ATGGCTTCCA  |
|     | GACTACGGCG | ACCGCTAAGT | CCAAGTAGTA | CGGCAAAACAC | TACCGAAGGT  |
| 701 | TGTCGGCAGA | ATGCTTAATG | AATTACAACA | GTACTGCCAT  | GAGTGGCAGG  |
|     | ACAGCCGTCT | TACGAATTAC | TTAATGTTGT | CATGACGCTA  | CTCACCGTCC  |
| 751 | GCGGGGCGTA | ATTTTTTTAA | GGCAGTTATT | GGGTGCCCTT  | AAACGCCTGG  |

Figure 34: functional map and sequence of pCAL module M17 (continued)

|     |             |           |            |            |            |
|-----|-------------|-----------|------------|------------|------------|
|     | CGCCCCCGCAT | TAAAAAATT | CCGTCAATAA | CCCACGGGAA | TTTGCGGACC |
|     |             | BglII     |            |            |            |
|     |             | ~~~~~     |            |            |            |
| 801 | TGCTAGATCT  | TCC       |            |            |            |
|     | ACGATCTAGA  | AGG       |            |            |            |

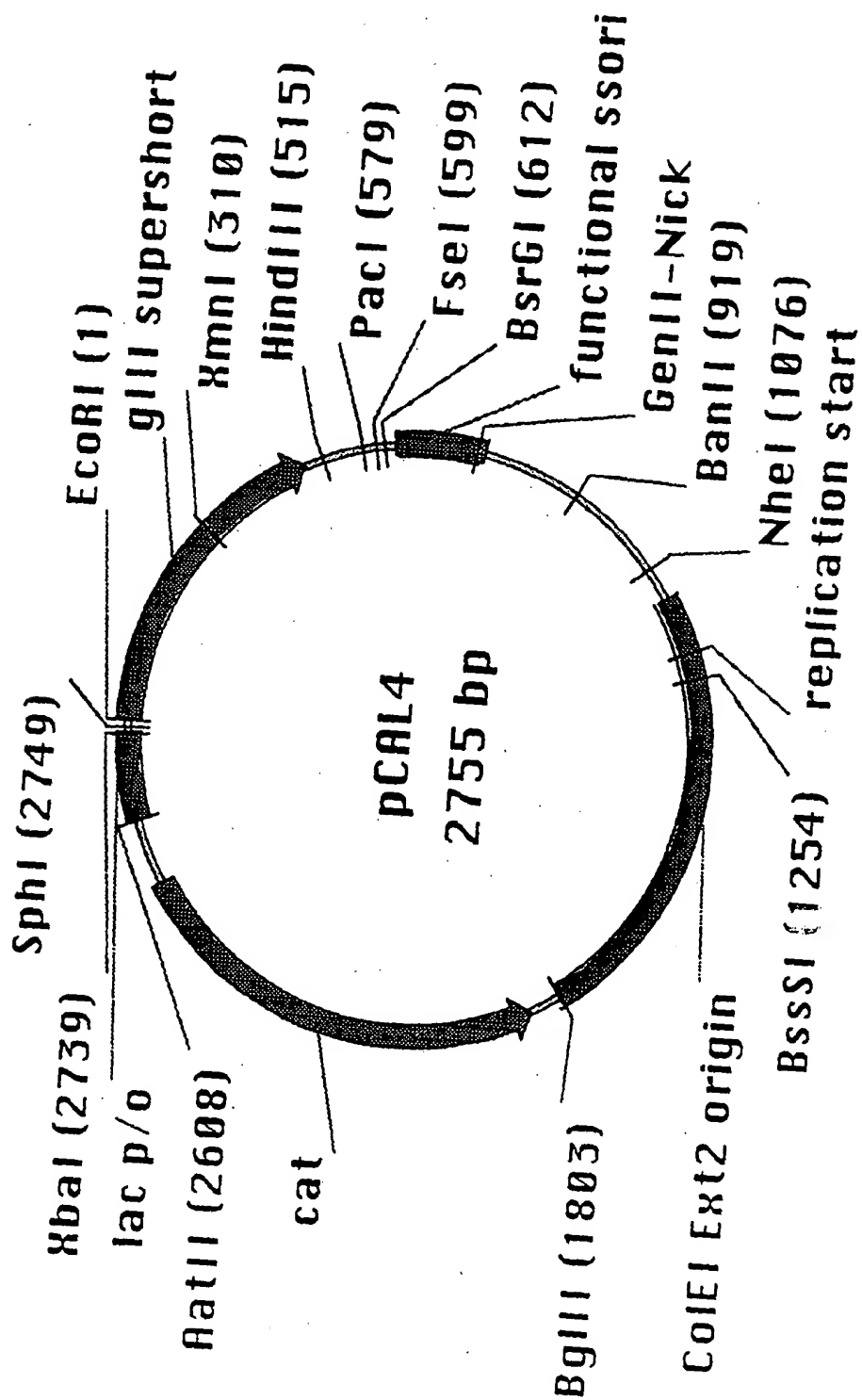


Figure 35: functional map and sequence of modular vector pCAL4



Figure 35: functional map and sequence of modular vector pCAL4 (continued)

## EcoRI

```

1  AATTCGAGCA GAAGCTGATC TCTGAGGAGG ATCTGTAGGG TGGTGGCTCT
   TTAAGCTCGT CTTCGACTAG AGACTCCTCC TAGACATCCC ACCACCGAGA

51  GGTTCGGTG ATTTTGATTA TGAAAAGATG GCAAACGCTA ATAAGGGGGC
   CCAAGGCCAC TAAAACTAAT ACTTTCTAC CGTTGCGAT TATCCCCCG

101 TATGACCGAA AATGCCGATG AAAACGCGCT ACAGTCTGAC GCTAAAGGCA
   ATACTGGCTT TTACGGCTAC TTTTGCGCGA TGTGAGACTG CGATTTCGCT

151 AACTTGATTC TGTCGCTACT GATTACGGTG CTGCTATCGA TGGTTTCATT
   TTGAACTAAG ACAGCGATGA CTAATGCCAC GACGATAGCT ACCAAAAGTAA

201 GGTGACGTTT CCGGCCCTTGC TAATGGTAAT GGTGCTACTG GTGATTTTGC
   CCACTGCAAA GGCCGGAACG ATTACCATTG CCACGATGAC CACTAAAACG

251 TGGCTCTAAT TCCCAAATGG CTCAAGTCGG TGACGGTGAT AATCACCTT
   ACCGAGATTA AGGTTTACC GAGTTCAGCC ACTGCCACTA TTAAGTGGA

```

## XmnI

```

~~~~~
301 TAATGAATAA TTCCCGTCAA TATTACCTT CCTCCCTCA ATCGGTTGAA
 ATTACTTATT AAAGGCAGTT ATAAATGGAA GGGAGGGAGT TAGCCAACTT

```

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

```

351 TGTCGCCCTT TTGTCTTTGG CGCTGGTAAA CCATATGAAT TTCTATTGA
 ACAGCGGGAA AACAGAAACC GCGACCATTT GTATACTTA AAAGATAACT

401 TTGTGACAAA ATAAACTTAT TCCGTGGTGT CTTTGCGTTT CTTTATATG
 AACACTGTTT TATTGAATA AGGCACCACA GAAACGCAA GAAAATATAC

451 TTGCCACCTT TATGTATGTA TTTTCTACGT TTGCTAACAT ACTGCGTAAT
 AACGGTGGAA ATACATACAT AAAAGATGCA AACGATTGTA TGACGCATTA

 HindIII
                                ~~~~~

501  AAGGAGTCTT GATAAGCTTG ACCTGTGAAG TGAAAAATGG CGCAGATTGT
    TTCCTCAGAA CTATTCGAAC TGGACACTTC ACTTTTACC GCGTCTAACA

                                PacI
                                ~~~~~

551 GCGACATTTT TTTTGTCTGC CGTTTAATTA AAGGGGGGG GGGCGCGCC
 CGCTGTAAAA AAACACAGACG GCAAATTAAT TTCCCCCCC CCGCGCCGG

 BsrGI
                                ~~~~~

601  TGGGGGGGGG TGTACATGAA ATTGTAACCG TTAATATTTT GTTAAATTC
    ACCCCCCCCC ACATGTACTT TAACATTTCG AATTATAAAA CAATTTTAAAG

```

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

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651  GCGTTAAATT TTTGTTAAAT CAGTCATTT TTTAAACCAAT AGGCCGAAAT
    CGCAATTAA AAACAATTAA GTCAGATAA AAATTGGTTA TCCGGCTTTA

701  CGGCAAAATC CCTATAAAT CAAAGAATA GACCGAGATA GGGTTGAGTG
    GCCGTTTATG GGAATATTAA GTTTTCTTAT CTGGCTCTAT CCCAACTCAC

751  TTGTTCCAGT TTGGAACAAG AGTCCACTAT TAAAGAACGT GGACTCCAAC
    AACAGGTCA AACCTTGTT TCAGGTGATA ATTTCTTGCA CCTGAGGTTG

801  GTCAAAGGC GAAAAACCGT CTATCAGGC GATGGCCAC TACGAGAACC
    CAGTTTCCCG CTTTTTGGCA GATAGTCCCG CTACCGGGTG ATGCTCTTGG

851  ATCACCCCTAA TCAAGTTT TTGGGTCGAG GTGCCGTAAA GCACTAAATC
    TAGTGGGATT AGTTCAAAAA ACCCCAGCTC CACGGCATTT CGTGATTAG

                                BanII
                                ~~~~~

901 GGAACCCCTAA AGGAGCCCC CGATTAGAG CTTGACGGG AAAGCCGGCG
 CCTTGGGATT TCCCTCGGG GCTAAATCTC GAACTGCCCC TTTCGGCCGC

951 AACGTGGCGA GAAAGGAAG GAAGAAAGCG AAAGAGCGG GCGCTAGGCG
 TTGCACCGCT CTTTCCCTTCC CTTCTTTCCG TTTCCTCGCC CGCGATCCCG

```

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|      |                                                         |       |
|------|---------------------------------------------------------|-------|
| 1001 | GCTGGCAAGT GTAGGGGTCA CGTGCGCGT AACCAACACA CCGCGCGCGC   |       |
|      | CGACCGTTCA CATCGCCAGT GCGACGCGCA TTGGTGGTGT GCGCGCGCGC  |       |
|      |                                                         |       |
|      |                                                         | NheI  |
|      |                                                         | ~~~~~ |
| 1051 | TTAATGCGCC GCTACAGGC GCGTGCTAGC CATGTGAGCA AAAGGCCAGC   |       |
|      | AATTACGCGG CGATGTCCCG CGCACGATCG GTACACTCGT TTTCGGGTCCG |       |
|      |                                                         |       |
| 1101 | AAAAGGCCAG GAACCGTAA AAGGCCGCGT TGCTGGCGTT TTCCCATAGG   |       |
|      | TTTTCGGTC CTTGGCATT TTCCGGCGCA ACGACCGCAA AAAGGTATCC    |       |
|      |                                                         |       |
| 1151 | CTCCGCCCCC CTGACGAGCA TCACAAAAT CGACGCTCAA GTCAGAGGTG   |       |
|      | GAGCGGGG GACTGCTCGT AGTGTTTTTA GCTGCGAGTT CAGTCTCCAC    |       |
|      |                                                         |       |
| 1201 | GCGAAACCCG ACAGGACTAT AAAGATACCA GCGGTTTCCC CCTGGAAGCT  |       |
|      | CGCTTTGGGC TGTCCTGATA TTTCATATGGT CCGCAAAGGG GGACCTTCGA |       |
|      |                                                         |       |
|      |                                                         | BssSI |
|      |                                                         | ~~~~~ |
| 1251 | CCCTCGTGCG CTCTCCTGTT CCGACCCCTGC CGCTTACCGG ATACCTGTCC |       |
|      | GGAGCACGC GAGAGGACAA GGCTGGGACG CGGAATGGCC TATGGACAGG   |       |
|      |                                                         |       |
| 1301 | GCCTTCTCC CTTCGGGAG CGTGGCGCTT TCTCATAGCT CACGCTGTAG    |       |
|      | CGGAAAGAGG GAAGCCCTTC GCACCGCGAA AGAGTATCGA GTGCCGACATC |       |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|      |             |             |             |             |            |
|------|-------------|-------------|-------------|-------------|------------|
| 1351 | GTATCTCAGT  | TCGGTGTAGG  | TCGTTTCGCTC | CAAGCTGGGC  | TGTGTGCACG |
|      | CATAGAGTCA  | AGCCACATCC  | AGCAAGCGAG  | GTTTCGACCCG | ACACACGTGC |
| 1401 | AACCCCCCGT  | TCAGCCCCGAC | CGCTGCCGCTT | TATCCGGGTAA | CTATCGTCTT |
|      | TTGGGGGGCA  | AGTCGGGCTG  | GCGACGCGGA  | ATAGGCCATT  | GATAGCAGAA |
| 1451 | GAGTCCAACC  | CGGTAAGACA  | CGACTTATCG  | CCACTGGCAG  | CAGCCACTGG |
|      | CTCAGGTTGG  | GCCATTCTGT  | GCTGAATAGC  | GGTGACCGTC  | GTCGGTGACC |
| 1501 | TAACAGGATT  | AGCAGAGCGA  | GGTATGTAGG  | CGGTGCTACA  | GAGTTCCTGA |
|      | ATTGTCCTAA  | TCGTCTCGCT  | CCATACATCC  | GCCACGATGT  | CTCAAGAACT |
| 1551 | AGTGGTGGCC  | TAACTACGGC  | TACACTAGAA  | GAACAGTATT  | TGGTATCTGC |
|      | TCACCAACCGG | ATTGATGCCG  | ATGTGATCTT  | CTTGTCATAA  | ACCATAGACG |
| 1601 | GCTCTGCTGT  | AGCCAGTTAC  | CTTCGGAAAA  | AGAGTTGGTA  | GCTCTTGATC |
|      | CGAGACGACA  | TCGGTCAATG  | GAAGCCTTTT  | TCTCAACCAT  | CGAGAACTAG |
| 1651 | CGGCAAAACAA | ACCACCGCTG  | GTAGCGGTGG  | TTTTTTTGTG  | TGCAAGCAGC |
|      | GCCGTTTGTT  | TGGTGGCGAC  | CATCGCCACC  | AAAAAAACAA  | ACGTTCTGTC |
| 1701 | AGATTACGCG  | CAGAAAAAAA  | GGATCTCAAG  | AAGATCCTTT  | GATCTTTTCT |
|      | TCTAATGCGC  | GTCTTTTTTT  | CCTAGAGTTC  | TTCTAGGAAA  | CTAGAAAAAG |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

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1751 ACGGGGTCTG ACGCTCAGTG GAACGAAAC TCACGTTAAG GGATTTGGT
 TGCCCCAGAC TCGGAGTCAC CTGCTTTTG AGTGCAATTC CCTAAACCA

 BglII
      ~~~~~
1801  CAGATCTAGC  ACCAGGCGTT  TAAGGCACC  AATAACTGCC  TTAAAAAAT
      GTCTAGATCG  TGGTCCGCAA  ATTCCCGTGG  TTATTGACGG  AATTTTTTA

1851  TACGCCCCGC  CCTGCCACTC  ATCGCAGTAC  TGTGTGAATT  CATTAAGCAT
      ATGCGGGGCG  GGACGGTGAG  TAGCGTCATG  ACAACATTAA  GTAATTCCGTA

1901  TCTGCCGACA  TGGAAGCCAT  CACAAACGGC  ATGATGAACC  TGAATCGCCA
      AGACGGCTGT  ACCTTCGGTA  GTGTTTGCCG  TACTACTTGG  ACTTAGCGGT

1951  GCGGCATCAG  CACCTTGTCG  CCTGCGGTAT  AATATTTGCC  CATAGTGAAA
      CGCCGTAGTC  GTGGAACAGC  GGAACGCATA  TTATAAACGG  GTATCACTTT

2001  ACGGGGGCGA  AGAAGTTGTC  CATATTGGCT  ACGTTTAAAT  CAAAACTGGT
      TGCCCCCGCT  TCTTCAACAG  GTATAACCGA  TGCAAAATTA  GTTTTGACCA

2051  GAAACTCACC  CAGGGATTGG  CTGAGACGAA  AAACATATTC  TCAATAAAC
      CTTTGAGTGG  GTCCCTAACC  GACTCTGCTT  TTTGTATAAG  AGTTATTGG

```

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|      |                                                         |                                 |
|------|---------------------------------------------------------|---------------------------------|
| 2101 | CTTTAGGGAA ATAGGCCAGG TTTTCACCGT AACACGCCAC ATCTTGCGGAA | TATCGGTCG TGGTATTCAC TCCAGAGCGA |
|      | GAAATCCCTT TATCCGGTCC AAAAGTGGCA TTGTGCCGTG TAGAACGCTT  | AGGTCTCGCT                      |
| 2151 | TATATGTGTA GAAACTGCCG GAAATCGTCG                        | ACCATAAGTG                      |
|      | ATATACACAT CTTTGACGGC CTTTAGCAGC                        | AGGTCTCGCT                      |
| 2201 | TGAAAACGTT TCAGTTTGCT CATGGAAAAC GGTGTAACAA GGTGAACAC   | CCACATTGTT                      |
|      | ACTTTTGCAA AGTCAAACGA GTACCTTTTG                        | CCCACATTGTG                     |
| 2251 | TATCCCATAT CACCAGCTCA CCGTCTTTCA TTGCCATACG GAACCTCCGGG | CTTGAGGCCCC                     |
|      | ATAGGGTATA GTGGTCGAGT GGCAGAAAGT AACGGTATGC             |                                 |
| 2301 | TGAGCATTCA TCAGGCGGGC AAGAATGTGA ATAAAGGCCG GATAAACTT   | CTATTTTGAA                      |
|      | ACTCGTAAGT AGTCCGCCCG TTTCTTACACT                       |                                 |
| 2351 | GTGCTTATTT TTCTTTACGG TCTTTAAAAA GGCCGTAATA TCCAGCTGAA  | AGGTCGACTT                      |
|      | CACGAATAAA AAGAAATGCC AGAAATTTT                         |                                 |
| 2401 | CGGTCTGGTT ATAGGTACAT TGAGCAACTG ACTGAAATGC CTCAAAATGT  | GAGTTTACAC                      |
|      | GCCAGACCAA TATCCATGTA ACTCGTTGAC                        |                                 |
| 2451 | TCTTTACGAT GCCATTGGGA TATATCAACG GTGGTATATC CAGTGATTTT  | GTCACATAAA                      |
|      | AGAAATGCTA CGGTAAACCCT ATATAGTTGC                       |                                 |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

```

2501  TTTCTCCATT TTAGCTTCCT TAGCTCCTGA AAATCTCGAT AACTCAAAAA
      AAAGAGGTAA AATCGAAGGA ATCGAGGACT TTTAGAGCTA TTGAGTTT
2551  ATACGCCCGG TAGTGATCTT ATTTCAATTAT GGTGAAAGTT GGAACCTCAC
      TATGCGGGCC ATCACTAGAA TAAAGTAATA CCACTTTCAA CCTGGAGTG

      AatII
      ~~~~~
2601 CCGACGTCTA ATGTGAGTTA GCTCACTCAT TAGGCACCCC AGGCTTTACA
 GGCTGCAGAT TACACTCAAT CGAGTGAGTA ATCCGTGGGG TCCGAAATGT
2651 CTTTATGCTT CCGGCTCGTA TGTGTGTGG AATTGTGAGC GGATAACAAT
 GAAATACGAA GGCCGAGCAT ACAACACACC TTAACACTCG CTAATTGTTA

 XbaI SphI
      ~~~~~
2701  TTCACACAGG AACAGCTAT GACCATGATT ACGAATTCT AGAGCATGCG
      AAGTGTGTCC TTTGTGCGATA CTGGTACTAA TGCTTAAAGA TCTCGTACGC

      EcoRI

2751  GGGG
      CCCC

```



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors

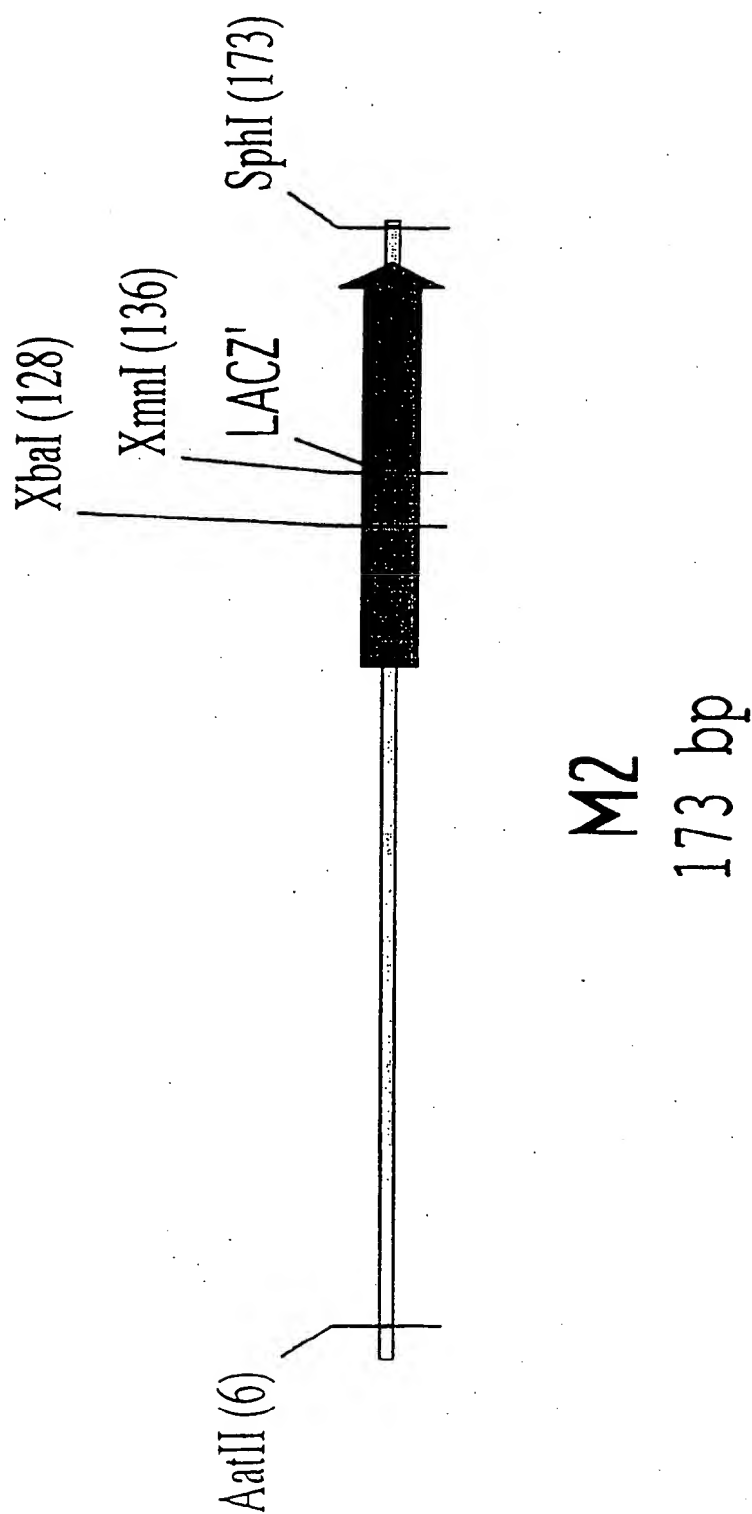


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 2:

AatII

~~~~~

1 GACGTCTTAA TGTGAGTTAG CTCACTCATT AGGCACCCCA GGCTTTACAC
CTGCAGAATT ACACTCAATC GAGTGAGTAA TCCGTGGGT CCGAAATGTG

51 TTTATGCTTC CGGCTCGTAT GTTGTGTGGA ATTGTGAGCG GATAACAATT
AAATACGAAG GCCGAGCATA CAACACACCT TAACACTCGC CTATTGTTAA

XmnI

~~~~~

XbaI

~~~~~

101 TCACACAGGA AACAGCTATG ACCATGTCTA GAATAACTTC GTATAATGTA
AGTGTGTCCT TTGTCGATAC TGGTACAGAT CTTATTGAAG CATATTACAT

SphI

~~~~~

151 CGCTATACGA AGTTATCGCA TGC  
GCGATATGCT TCAATAGCGT ACG

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

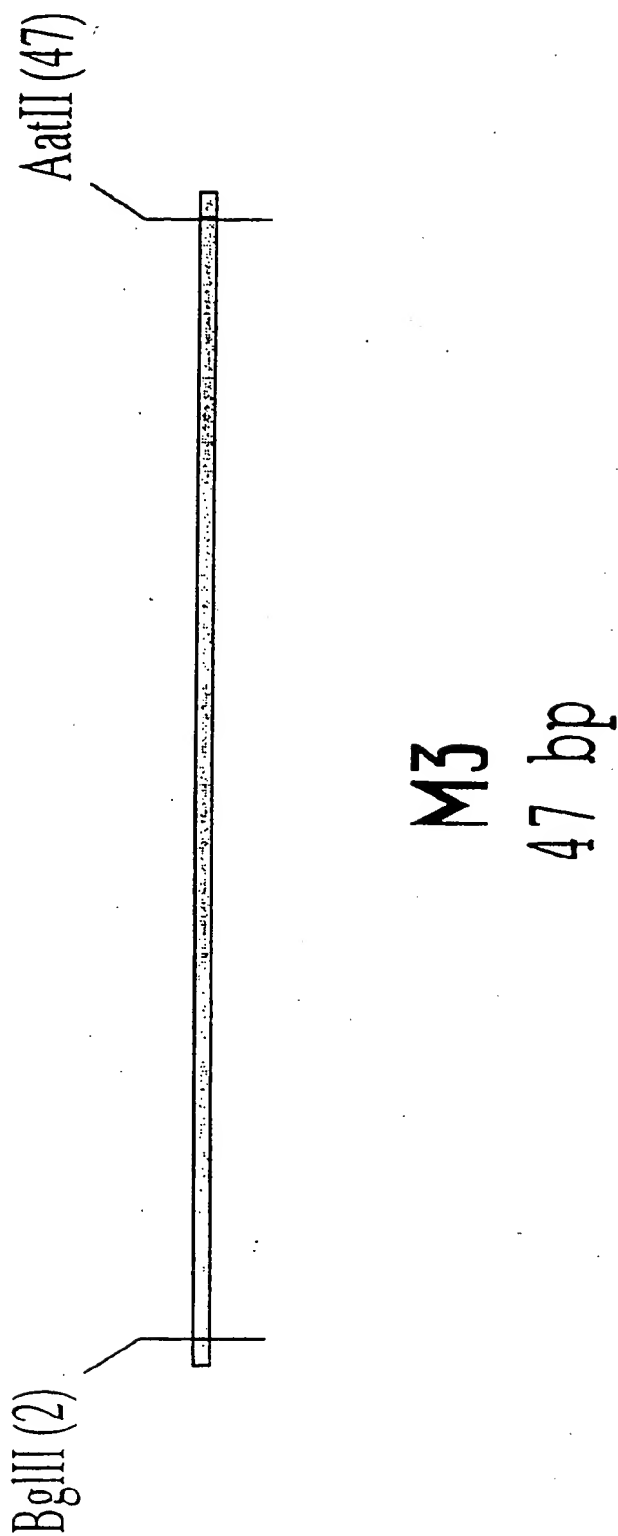


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 3:

|   | BglII                                               | AatII |
|---|-----------------------------------------------------|-------|
|   | -----                                               | ----- |
| 1 | AGATCTCATA ACTTCGTATA ATGTATGCTA TACGAAGTTA TGACGTC |       |
|   | TCTAGAGTAT TGAAGCATAT TACATACGAT ATGCTTCAAT ACTGCAG |       |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

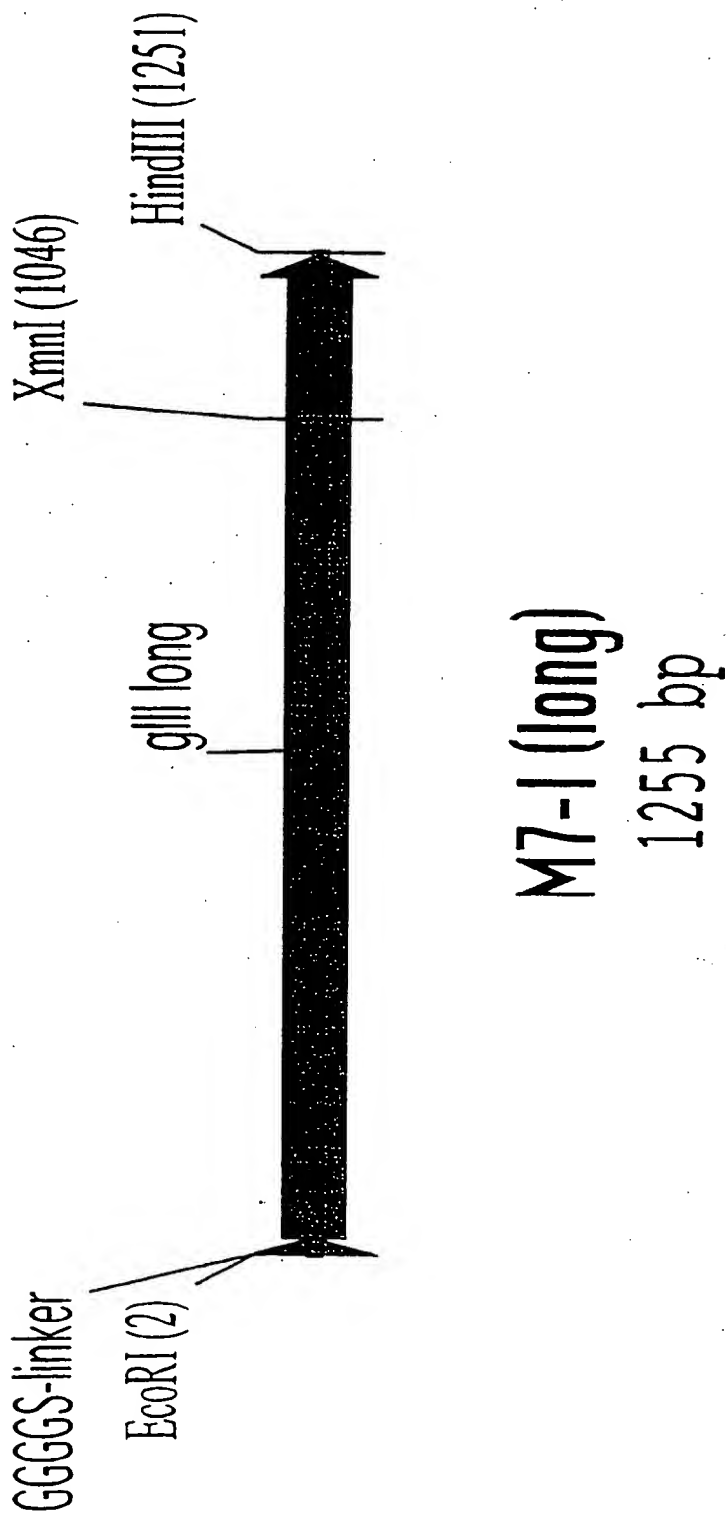


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

## M 7-I (long):

## ECORI

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-----  
1  GAATTCGGTG GTGGTGGATC TGCGTGGGCT GAAACGGGTG AAAGTTGTTT  
   CTTAAGCCAC CACCACCTAG ACGCACGCCA CTTTGCCAAC TTCAACAAA  
  
51  AGCAAAATCC CATAAGAAA ATTCATTAC TAACGCTCTG AAAGACGACA  
   TCGTTTTAGG GTATGTCTTT TAAGTAAATG ATTGCAGACC TTCTGCTGT  
  
101 AACTTTAGA TCGTTACGCT AACTATGAGG GCTGTCTGTG GAATGCTACA  
   TTTGAAATCT AGCAATGCCA TTGATACTCC CGACAGACAC CTTACGATGT  
  
151 GCGGTTGTAG TTTGTACTGG TGACGAAACT CAGTGTTACG GTACATGGGT  
   CCGCAACATC AACATGACC ACTGCTTTGA GTCACAATGC CATGTACCCA  
  
201 TCCTATTGGG CTTGCTATCC CTGAAAATGA GGGTGGTGGC TCTGAGGGTG  
   AGGATAACCC GAACGATAGG GACTTTTACT CCCACCACCG AGACTCCCAC  
  
251 GCGGTTCTGA GGGTGGCGGT TCTGAGGGTG GCGGTACTAA ACCTCCTGAG  
   CGCCAAGACT CCCACCGCCA AGACTCCCCAC CGCCATGATT TGGAGGACTC  
  
301 TACGGTGATA CACCTATTCC GGGCTATACT TATATCAACC CTCTCGACGG  
   ATGCCACTAT GTGGATAAGG CCCGATATGA ATATAGTTGG GAGAGCTGCC
```

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |                                                                                                                  |
|-----|------------------------------------------------------------------------------------------------------------------|
| 351 | CACTATACCG CCTGGTACTG AGCAAAACCC CGCTAATCCT AATCCTTCTC<br>GTGAATAGGC GGACCATGAC TCGTTTGGG GCGATTAGGA TTAGGAAGAG  |
| 401 | TTGAGGAGTC TCAGCCTCTT AATACTTTCA TGTTTCAGAA TAATAGGTTC<br>AACTCCTCAG AGTCGGAGAA TTATGAAAGT ACAAAGTCTT ATTATCCAAG |
| 451 | CGAAATAGGC AGGGGCATT AACTGTTTAT ACGGGCACTG TTAACAAGG<br>GCTTTATCCG TCCCCCGTAA TTGACAAATA TGCCCCGTGAC AATGAGTTCC  |
| 501 | CACTGACCCG GTTAAACTT ATTACCAGTA CACTCCTGTA TCATCAAAAG<br>GTGACTGGGG CAATTTTGAA TAATGGTCAT GTGAGGACAT AGTAGTTTC   |
| 551 | CCATGTATGA CGCTTACTGG AACGGTAAAT TCAGAGACTG CGCTTTCCAT<br>GGTACATACT GCGAATGACC TTGCCATTTA AGTCTCTGAC GCGAAAGGTA |
| 601 | TCTGGCTTTA ATGAGGATTT ATTTGTTTGT GAATATCAAG GCCAATCGTC<br>AGACCGAAAT TACTCCTAAA TAAACAAACA CTTATAGTTC CGGTTAGCAG |
| 651 | TGACCTGCCT CAACCTCCTG TCAATGCTGG CGGCGGCTCT GTGGTGGTT<br>ACTGGACGGA GTTGAGGAC AGTTACGACC GCCGCCGAGA CCACCACCAA   |
| 701 | CTGGTGGCGG CTCTGAGGGT GGTGGCTCTG AGGGTGGCGG TTCTGAGGGT<br>GACCACCGCC GAGACTCCCA CCACCGAGAC TCCCACCGCC AAGACTCCCA |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |            |            |             |             |            |
|------|------------|------------|-------------|-------------|------------|
| 751  | GGCGGCTCTG | AGGGAGGCGG | TTCCGGTGGT  | GGCTCTGGTT  | CCGGTGATTT |
|      | CCGCCGAGAC | TCCCTCCGCC | AAGGCCACCA  | CCGAGACCAA  | GGCCACTAAA |
| 801  | TGATTATGAA | AAGATGGCAA | ACGCTAAATA  | GGGGGCTATG  | ACCGAAAATG |
|      | ACTAATACTT | TTCTACCGTT | TGCGATTATT  | CCCCCGATAC  | TGGCTTTTAC |
| 851  | CCGATGAAAA | CGCGCTCAG  | TCTGACGCTA  | AAGGCCAACT  | TGATTCTGTC |
|      | GGCTACTTTT | CGCGATGTC  | AGACTGCCGAT | TTCCGTTTGA  | ACTAAGACAG |
| 901  | GCTACTGATT | ACGGTGCTGC | TATCGATGGT  | TTCATTGGTG  | ACGTTTCCGG |
|      | CGATGACTAA | TGCCACGACG | ATAGCTACCA  | AAGTAACCAC  | TGCAAAGGCC |
| 951  | CCTTGCTAAT | GGTAATGGTG | CTACTGGTGA  | TTTTTGCTGGC | TCTAATTCCC |
|      | GGAACGATTA | CCATTACCAC | GATGACCACT  | AAAACGACCG  | AGATTAAAGG |
| XmnI |            |            |             |             |            |
| 1001 | AAATGGCTCA | AGTCGGTGAA | GGTGATAATT  | CACCTTTAAT  | GAATAATTTC |
|      | TTTACCGAGT | TCAGCCACTT | CCACTATTAA  | GTGGAAATTA  | CTTATTAAAG |
| 1051 | CGTCAATATT | TACCTTCCAT | CCCTCAATCG  | GTTGAATGTC  | GCCCTTTTGT |
|      | GCAGTTATAA | ATGGAAGGTA | GGGAGTTAGC  | CAACTTACAG  | CGGGAACAAC |



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |            |             |            |            |            |
|------|------------|-------------|------------|------------|------------|
| 1101 | CTTTGGCGCT | GGTAAACCCCT | ATGAATTTC  | TATTGATTGT | GACAAAATAA |
|      | GAAACCGCGA | CCATTGGGA   | TACTTAAAG  | ATAACTAACA | CTGTTTATT  |
| 1151 | ACTTATTCCG | TGGTGTCCTT  | GCGTTTCTTT | TATATGTTGC | CACCTTTATG |
|      | TGAATAAGGC | ACCACAGAAA  | CGCAAAGAAA | ATATACAACG | GTGGAAATAC |
|      |            |             |            |            | HindIII    |
| 1201 | TATGTATTTT | CTACGTTTGC  | TAACTACTG  | CGTAATAAGG | AGTCTTGATA |
|      | ATACATAAAA | GATGCAAACG  | ATTGTATGAC | GCATTATTCC | TCAGAACTAT |
|      |            |             |            |            | HindI      |
|      |            |             |            |            | ~~~~       |
| 1251 |            | AGCTT       |            |            |            |
|      |            | TCGAA       |            |            |            |

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Figure 35a: Functional maps and sequences of additional: pCAL vector modules and pCAL vectors (continued)

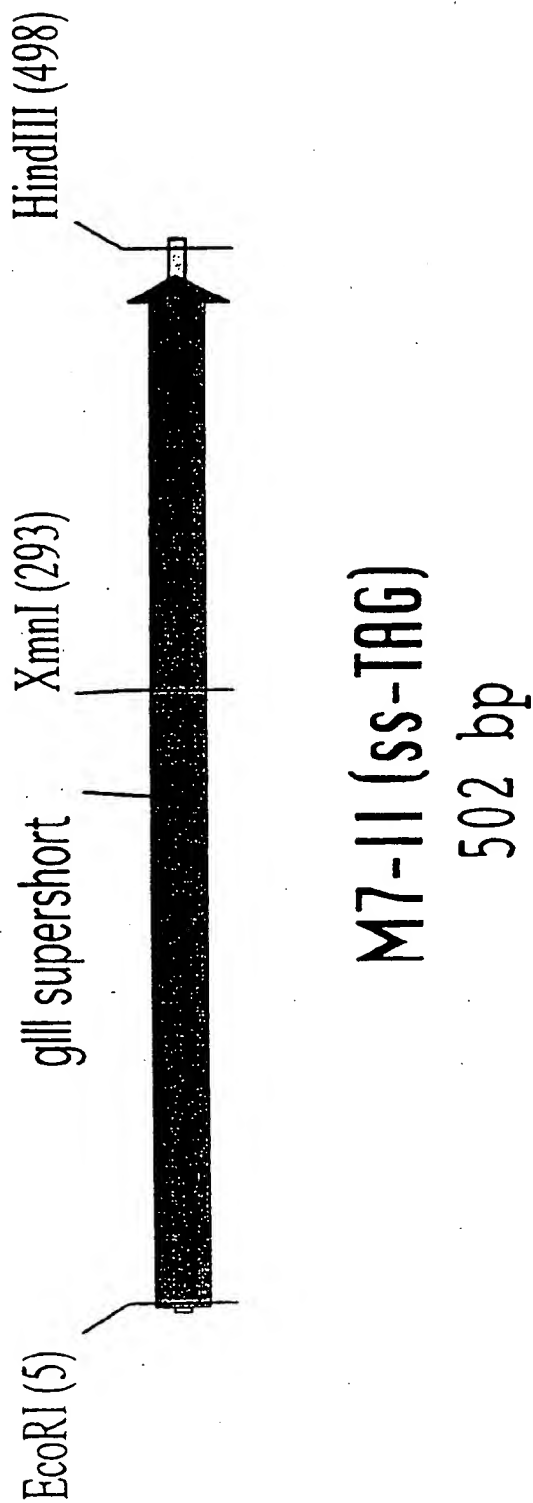


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

## M 7-II (ss-TAG) :

ECORI

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|     |             |            |             |            |            |
|-----|-------------|------------|-------------|------------|------------|
| 1   | CGGGAATTCTG | GAGCGGTTC  | CGGTGGTGGC  | TCTGGTTCCG | GTGATTTTGA |
|     | GCCCTTAAGC  | CTCCGCCAAG | GCCACCACCG  | AGACCAAGGC | CACTAAAACT |
| 51  | TTATGAAAAG  | ATGGCAAACG | CTAATAAGGG  | GGCTATGACC | GAAAATGCCG |
|     | AATACTTTC   | TACCGTTTGC | GATTATTCCC  | CCGATACTGG | CTTTTACGGC |
| 101 | ATGAAAACGC  | GCTACAGTCT | GACGCTAAAG  | GCAAACCTGA | TTCTGTCCGT |
|     | TACTTTTGCG  | CGATGTCAGA | CTGCGATTTC  | CGTTTGAAC  | AAGACAGCGA |
| 151 | ACTGATTACG  | GTGCTGCTAT | CGATGGTTTC  | ATTGGTGACG | TTTCCGGCCT |
|     | TGACTAATGC  | CACGACGATA | GCTACCAAAG  | TAACCACTGC | AAAGGCCCGA |
| 201 | TGCTAATGGT  | AATGGTGCTA | CTGGTGATT   | TGCTGGCTCT | AATCCCATAA |
|     | ACGATTACCA  | TTACCACGAT | GACCACTAAA  | ACGACCGAGA | TTAAGGGTTT |
| 251 | TGGCTCAAGT  | CGGTGACGGT | GATAATTCAC  | CTTTAATGAA | TAATTCCGT  |
|     | ACCGAGTTCA  | GCCACTGCCA | CTATTAAAGTG | GAAATTACTT | ATTAAAGGCA |

XmnI

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |                                                        |           |
|-----|--------------------------------------------------------|-----------|
| 301 | CAATATTAC CTTCCCTCCC TCAATCGGTT GAATGTCGCC CTTTGTCTT   | GAAACAGAA |
|     | GTTATAAATG GAAGGGAGGG AGTTAGCCAA CTTACAGCGG            |           |
| 351 | TGGCGCTGGT AAACCATATG AATTTCTTAT TGATTGTGAC AAAATAAACT |           |
|     | ACCGGACCA TTTGGTATAC TTAAAAGATA ACTAACACTG TTTTATTGA   |           |
| 401 | TATCCGTGG TGTCTTTCGG TTTCTTTTAT ATGTTGCCAC CTTTATGTAT  |           |
|     | ATAAGGCACC ACAGAAACGC AAAGAAAATA TACAACGGTG GAAATACATA |           |
|     |                                                        | HindIII   |
| 451 | GTATTTCTA CGTTGCTAA CATACTGCCGT AATAAGGAGT CTTGATAAGC  |           |
|     | CATAAAAGAT GCAAACGATT GTATGACGCA TTATTCCTCA GAACTATTCG |           |
| 501 | Hi                                                     | ---       |
|     | ~                                                      |           |
|     | TT                                                     |           |
|     | AA                                                     |           |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

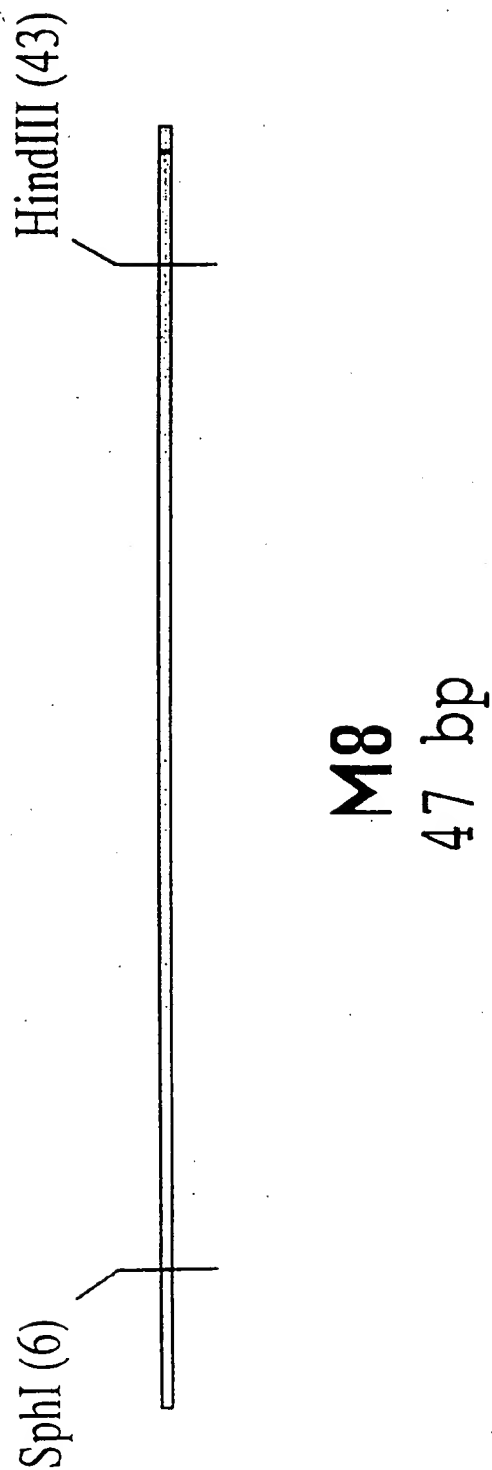


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 8:

|   |                                                     |  |         |
|---|-----------------------------------------------------|--|---------|
|   | SphI                                                |  | HindIII |
|   | ~~~~~                                               |  | ~~~~~   |
| 1 | GCATGCCATA ACTTCGTATA ATGTACGCTA TACGAAGTTA TAAGCTT |  |         |
|   | CGTACGGTAT TGAAGCATAT TACATGCGAT ATGCTTCAAT ATTCGAA |  |         |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

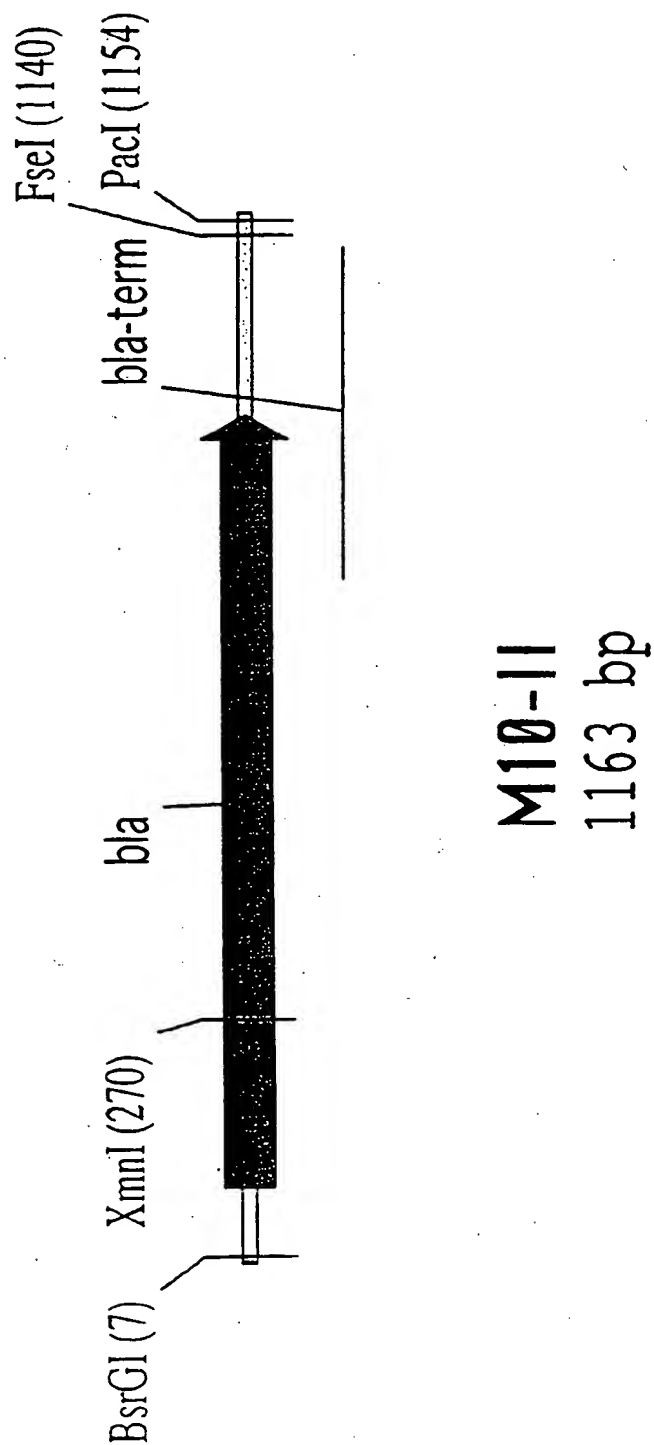


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

## M 10-II:

## BsrGI

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```

1  GGGGGTGTAC ATTCAAATAT GTATCCGCTC ATGAGACAAT AACCCGTGATA
   CCCCACATG TAAGTTTATA CATAGGCGAG TACTCTGTTA TTGGGACTAT

51  AATGCTTCAA TAATATTGAA AAAGGAAGAG TATGAGTATT CAACATTTC
   TTACGAAGTT ATTATAACTT TTTCCCTTCTC ATACTCATAA GTTGTAAGG

101 GTGTCGCCCT TATCCCTTT TTTGCGGCAT TTTGCCCTTCC TGTTTTTGCT
   CACAGCGGGA ATAAGGGAAA AAACGCCGTA AAACGGAAGG ACAAAAACGA

151 CACCCAGAAA CGCTGGTGAA AGTAAAAGAT GCTGAGGATC AGTTGGGTGC
   GTGGGTCTTT GCGACCACTT TCATTTTCTA CGACTCCTAG TCAACCCACG

201 GCGAGTGGGT TACATCGAAC TGGATCTCAA CAGCGGTAAG ATCCTTGAGA
   CGCTCACCCA ATGTAGCTTG ACCTAGAGTT GTCGCCATTC TAGGAACTCT

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XmnI

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251 GTTTTCGCCC CGAAGAACGT TTCCAATGA TGAGCACTTT TAAAGTTCTG
   CAAAAGCGGG GCTTCTTGCA AAAGTTACT ACTCGTGAAA ATTCAAGAC

```



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

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301  CTATGTGGCG  CCGTATTATC  CCGTATTGAC  GCCGGGCAAG  AGCAACTCGG
      GATACACCGC  GCCATAATAG  GGCATAACTG  CGGCCCGTTC  TCGTTGAGCC

351  TCGCCCGCATA  CACTATTCTC  AGAATGACTT  GGTGAGTAC   TCACCAGTCA
      AGCGGCGGTAT  GTGATAAGAG  TCTTACTGAA  CCAACTCATG  AGTGGTCAGT

401  CAGAAAAGCA  TCTTACGGAT  GGCATGACAG  TAAGAGAATT  ATGCAGTGCT
      GTCCTTTTCGT  AGAATGCCCTA  CCGTACTGTC  ATTCTCTTAA  TACGTCACGA

451  GCCATAACCA  TGAGTGATAA  CACTGCGGCC  AACTTACTTC  TGACAAACGAT
      CCGTATTGGT  ACTCACTATT  GTGACGCCGG  TTGAATGAAG  ACTGTTGCTA

501  CGGAGGACCG  AAGGAGCTAA  CCGCTTTTTC  GCACAACATG  GGGATCATG
      GCCTCCTGGC  TTCCTCGATT  GCGAATAAAA  CGTGTGTAC   CCCCTAGTAC

551  TAACTCGCCT  TGATCGTTGG  GAACCGGAGC  TGAATGAAGC  CATAACAAAC
      ATGAGCGGA  ACTAGCAACC  CTTGGCCTCG  ACTTACTTCG  GTATGGTTTG

601  GACGAGCGTG  ACACCACGAT  GCCTGTAGCA  ATGGCAACAA  CGTTGCGCAA
      CTGCTCGCAC  TGTGGTGCTA  CGGACATCGT  TACCGTTGTT  GCAACGCGTT

651  ACTATTAACT  GGCGAACTAC  TTA CTCTAGC  TTCCCGGCAA  CAGTTAATAG
      TGATAATTGA  CCGCTTGATG  AATGAGATCG  AAGGGCCGTT  GTCAATTATC

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |                                                         |                                                        |
|------|---------------------------------------------------------|--------------------------------------------------------|
| 701  | ACTGGATGGA GCGGGATAAA GTTGCAGGAC CACTTCTGCG CTCGGCCCCTT | TGACCTACCT CCGCCTATT CAACGTCCTG GTGAAGACGC GAGCCGGGAA  |
| 751  | CCGGCTGGCT GGTTATTGC TGATAAATCT GGAGCCGGTG AGCGTGGGTC   | GGCCGACCGA CCAAATAACG ACTATTAGA CCTCGGCCAC TCGCACCCAG  |
| 801  | TCGCGGTATC ATTGCAGCAC TGGGGCCAGA TGGTAAGCCC TCCC GTATCG | AGCGCCATAG TAACGTCGTG ACCCCGGTCT ACCATTCTGG AGGCGATAGC |
| 851  | TAGTTATCTA CACGACGGGG AGTCAGGCAA CTATGGATGA ACGAAATAGA  | ATCAATAGAT GTGCTGCCCC TCAGTCCGTT GATACCTACT TGCTTTATCT |
| 901  | CAGATCGCTG AGATAGGTGC CTCACTGATT AAGCATTGGG TAACTGTCAG  | GTCTAGCGAC TCTATCCACG GAGTGAATAA TTCGTAACCC ATTGACAGTC |
| 951  | ACCAAGTTA CTCATATATA CTTTAGATTG ATTTAAAACT TCATTTTAA    | TGGTTCAAAT GAGTATATAT GAAATCTAAC TAAATTTTGA AGTAAAAATT |
| 1001 | TTTAAAAGGA TCTAGGTGAA GATCCCTTTT GATAATCTCA TGACCAAAT   | AAATTTTCCT AGATCCACTT CTAGGAAAAA CTATTAGAGT ACTGGTTTA  |
| 1051 | CCCTTAACGT GAGTTTTCGT TCCACTGAGC GTCAGACCCC GTAGAAAAGA  | GGGAATTGCA CTCAAAAGCA AGGTGACTCG CAGTCTGGG CATCTTTCT   |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |                                                      | FseI  | PacI |
|------|------------------------------------------------------|-------|------|
|      |                                                      | ~~~~~ | ~~   |
| 1101 | TCAAAGGATC TTCTTGAGAT CCTTTTGAT AATGGCCGGC CCCCCCCTT |       |      |
|      | AGTTTCCTAG AAGAACTCTA GGAAAACTA TTACCGGCCG GGGGGGGAA |       |      |
|      | PacI                                                 |       |      |
|      | ~~~~~                                                |       |      |
| 1151 | AATTAAGGG GGG                                        |       |      |
|      | TTAATTCCCC CCC                                       |       |      |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

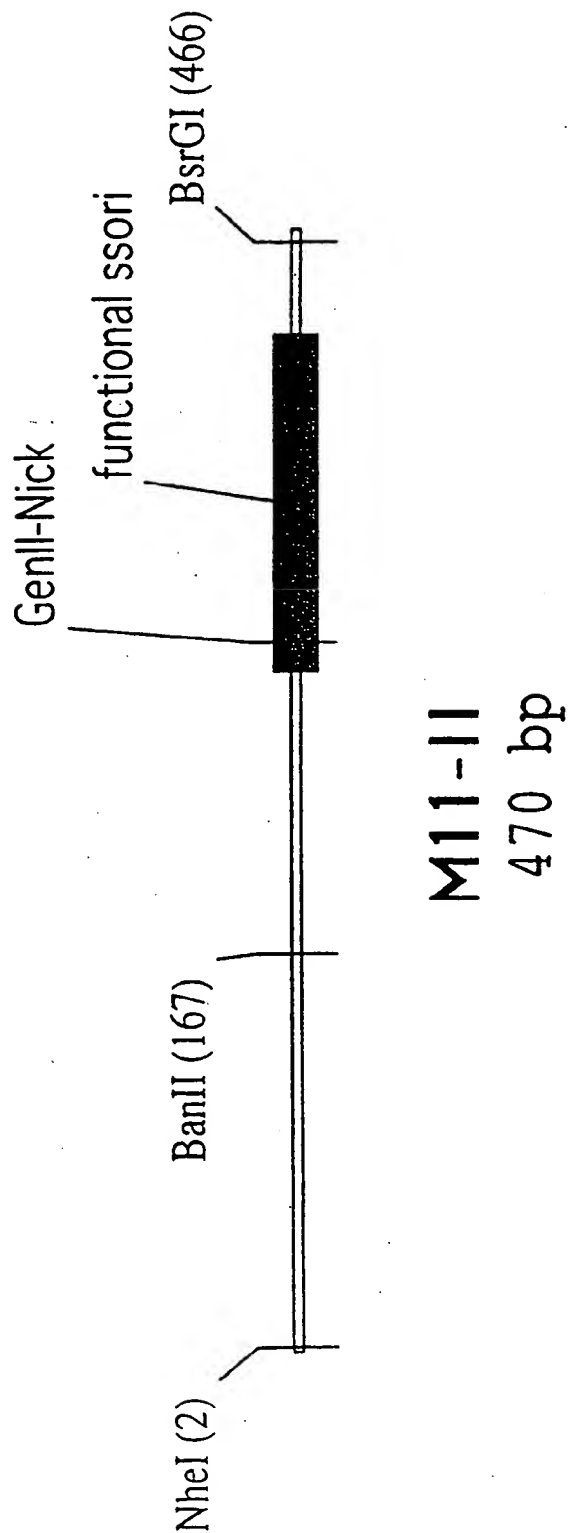


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

## M11-II:

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1 GCTAGCACGC GCCCTGTAGC GCGGCATTAA GCGCGGCGGG TGTGGTGGTT
CGATCGTGCG CCGGACATCG CCGCGTAATT CCGCGCGCCC ACACCACCAA

51 ACGCGCAGCG TGACCGCTAC ACTTGCCAGC GCCCTAGCGC CCGCTCCTTT
TGCGCGTCGC ACTGGCGATG TGAACGGTCG CCGGATCGCG GCGAGGAAA

101 CGCTTCTTC CCTTCCTTTC TCGCCACGTT CGCGGGCTTT CCCCGTCAAG
GCGAAAGAAG GGAAGGAAAG AGCGGTGCAA GCGGCCGAAA GGGCAGTTC

BanII

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151 CTCTAAATCG GGGCTCCCT TTAGGGTCC GATTAGTGC TTACGGCAC  
GAGATTAGC CCCGAGGGA AATCCCAAGG CTAATCACC AAATGCCGTC

201 CTCGACCCCA AAAAATTGA TTAGGGTGAT GGTTCTCGTA GTGGGCCATC  
GAGCTGGGT TTTTGAAC TATCCACTA CCAAGAGCAT CACCCGGTAG

251 GCCCTGATAG ACGGTTTTC GCCCTTTGAC GTTGGAGTCC ACGTTCTTTA  
CGGACTATC TGCCAAAAG CGGAAACTG CAACCTCAGG TGCAAGAAAT

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

```
301  ATAGTGGACT CTTGTTCCAA ACTGGAACAA CACTCAACCC TATCTCGGTC
    TATCACCTGA GAACAAGGTT TGACCTTGTT GTGAGTTGGG ATAGAGCCAG

351  TATTCTTTTG ATTTATAAGG GATTTGCCG ATTTCGGCCT ATTGGTTAAA
    ATAAGAAAC TAAATATTCC CTAAACCGG TAAAGCCGGA TAACCAATTT

401  AAATGAGCTG ATTTAACAAA AATTAAACGC GAATTTTAAC AAAATATTAA
    TTTACTCGAC TAAATTGTTT TTAAATTGCG CTTAAAAATTG TTTTATAATT

451  CGTTTACAAT TTCATGTACA
    GCAAATGTTA AAGTACATGT
```

BSrGI

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

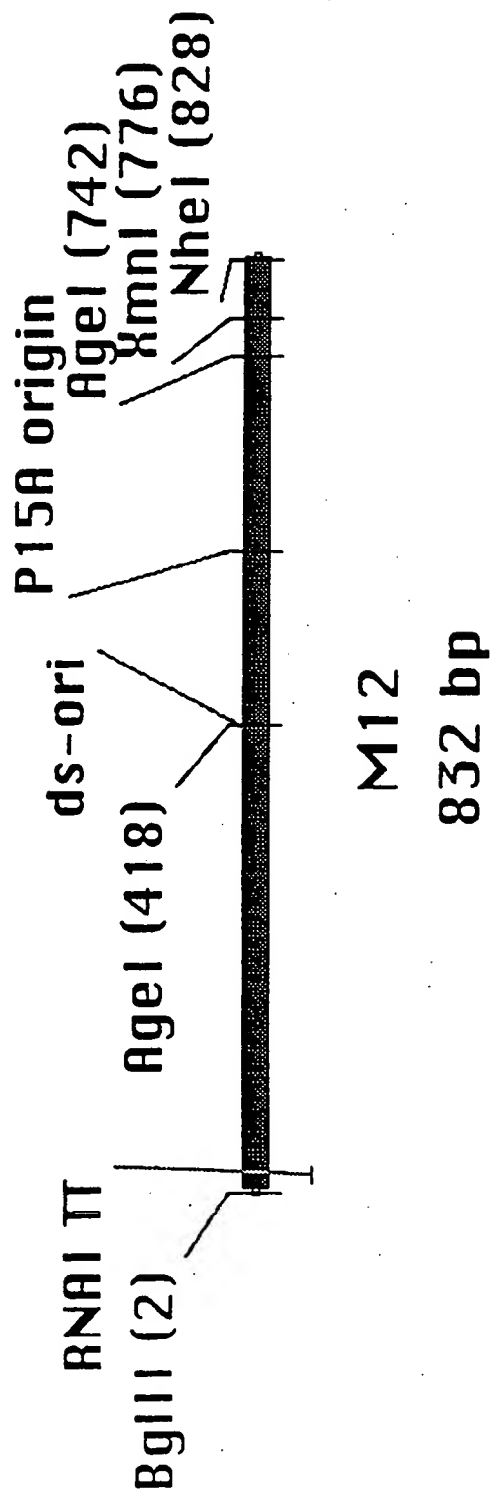


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 12:		BgIII	
		~~~~~	
1	AGATCTAATA	AGATGATCTT	CTTGAGATCG
	TCTAGATTAT	TCTACTAGAA	GAACTCTAGC
			AAAAACCAGAC
			GCGCATTAGA
51	CTTGCTCTGA	AAACGAAAAA	ACCGCCTTGC
	GAACGAGACT	TTTGCTTTTT	TGGCGGAACG
			TCCCGCCAAA
			AAGCATCCAA
			TTCTGTAGGTT
101	CTCTGAGCTA	CCAACTCTTT	GAACCGAGGT
	GAGACTCGAT	GGTTGAGAAA	CTTGGCTCCA
			TTGACCCGAAC
			CTCCTCGCGT
151	GTCACTAAAA	CTTGTCCTTT	CAGTTAGCC
	CAGTGATTTT	GAACAGGAAA	GTCAAATCGG
			AATTGGCCGC
			GTACTGAAGT
201	AGACTAACTC	CTCTAAATCA	ATTACCAGTG
	TCTGATTGAG	GAGATTTAGT	TAAATGGTCAC
			CGACGACGGT
			CACACGAAA
251	TGCATGTCTT	TCCGGGTTGG	ACTCAAGACG
	ACGTACAGAA	AGGCCCAACC	TGAGTTCTGC
			TATCAATGGC
			CTATTCCGCG
301	AGCGGTCGGA	CTGAACGGGG	GGTTCGTGCA
	TCGCCAGCCT	GACTTGCCCC	CCAAGCACGT
			ATGTCAGGTC
			GAACCTCGCT



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

351	ACTGCCCTACC	CGGAACCTGAG	TGTCAGGCGT	GGAATGAGAC	AAACGCGGCC
	TGACGGATGG	GCCTTGACTC	ACAGTCCGCA	CCTTACTCTG	TTTGCGGCCGG
AgeI					
~~~~~					
401	ATAACAGCGG	AATGACACCG	GTAACCCGAA	AGGCAGGAAC	AGGAGAGCGC
	TATTGTCGCC	TTACTGTGGC	CATTGGCTT	TCCGTCCCTTG	TCCTCTCGCG
451	AGGAGGGAGC	CGCCAGGGGG	AAACGCCCTGG	TATCTTTATA	GTCCTGTCTGG
	TCCTCCCTCG	GCGGTCCCCC	TTTGCGGACC	ATAGAAATAT	CAGGACAGCC
501	GTTTCGCCAC	CACTGATTG	AGCGTCAGAT	TTCGTGATGC	TTGTCAGGGG
	CAAAGCGGTG	GTGACTAAAC	TCGCAGTCTA	AAGCACTACG	AACAGTCCCC
551	GGCGGAGCCT	ATGGAAAAAC	GGCTTTGCCG	CGGCCCTCTC	ACTTCCCTGT
	CCGCCCTCGA	TACCTTTTG	CCGAAACGGC	GCCGGGAGAG	TGAAGGGACA
601	TAAGTATCTT	CCTGGCATCT	TCCAGGAAAT	CTCCGCCCCG	TTCGTAAGCC
	ATTCATAGAA	GGACCGTAGA	AGGTCCTTTA	GAGCGGGGC	AAGCATTCGG
651	ATTTCCGCTC	GCCGCAGTCG	AACGACCGAG	CGTAGCGAGT	CAGTGAGCGA
	TAAAGGCGAG	CGGCGTCAGC	TTGCTGGCTC	GCATCGCTCA	GTCACTCGCT

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

				AgeI	
				~~~~~	
701	GGAAGCGGAA	TATATCCTGT	ATCACATATT	CTGCTGACGC	ACCGGTGCAG
	CCTTCGCCCTT	ATATAGGACA	TAGTGTATAA	GACGACTGCG	TGGCCACGTC
			XmnI		
			~~~~~		
751	CCTTTTCTTCT	CCTGCCACAT	GAAGCACTTC	ACTGACACCC	TCATCAGTGC
	GGAAAAAAGA	GGACGGTGTA	CTTCGTGAAG	TGACTGTGGG	AGTAGTCACG
			NheI		
			~~~~~		
801	CAACATAGTA	AGCCAGTATA	CACTCCGCTA	GC	
	GTGTATCAT	TCGGTCATAT	GTGAGGCGAT	CG	

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

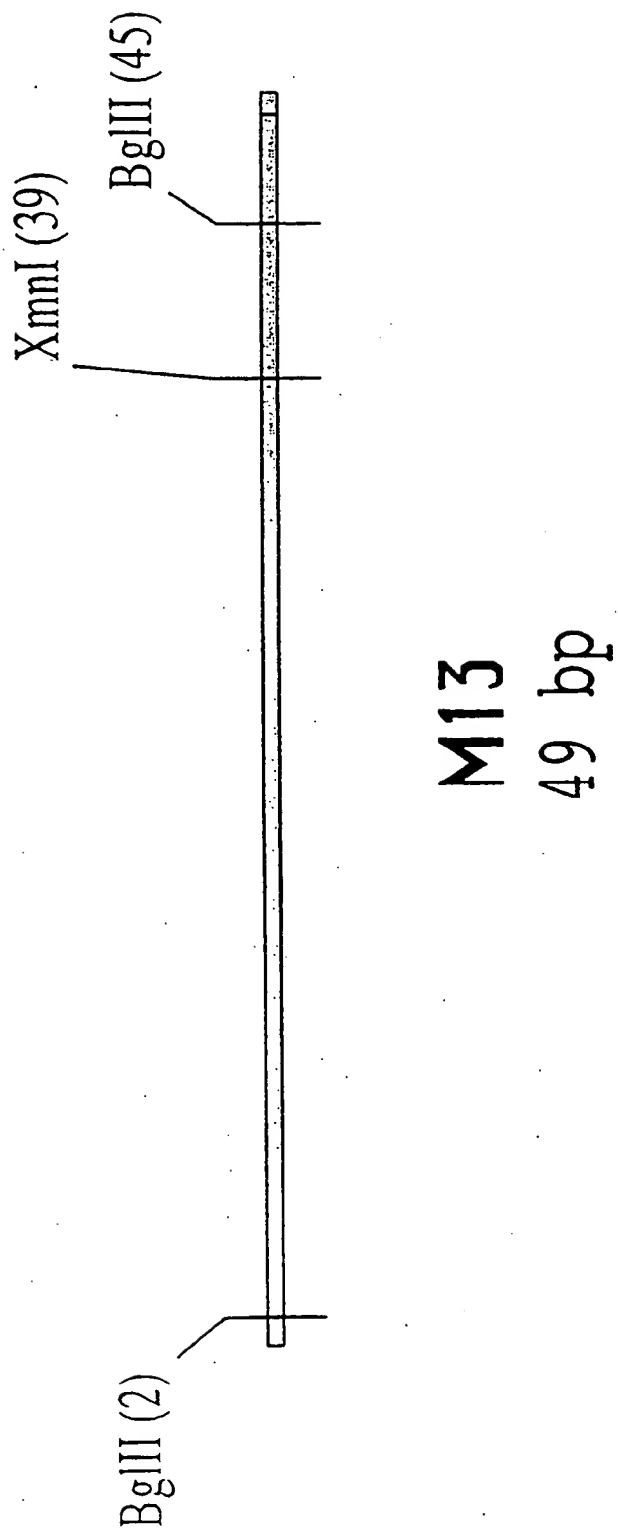


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 13:

	BglII	XmnI	BglII
	-----	-----	-----
1	AGATCTCATA	ATGTATGCTA	TACGAAGTTA
	TCTAGAGTAT	TGAAGCATAT	TACATACGAT
			ATGCTTCAAT
			AAGTCTAGA

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

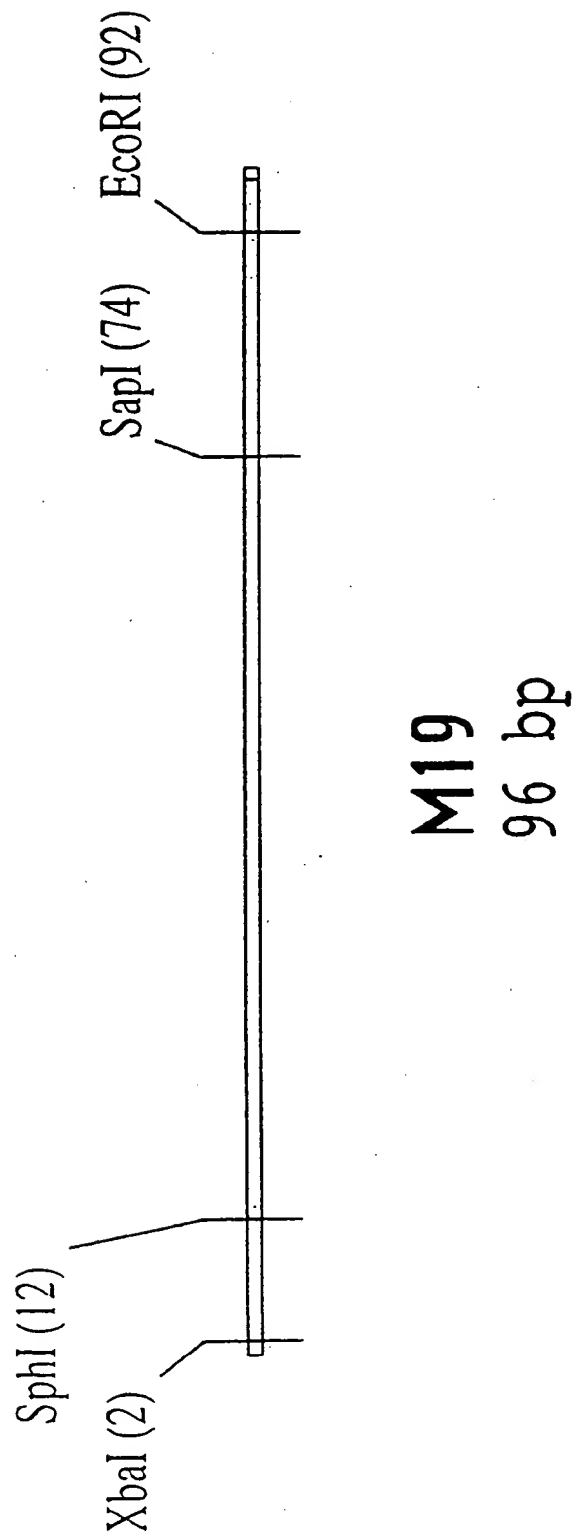




Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

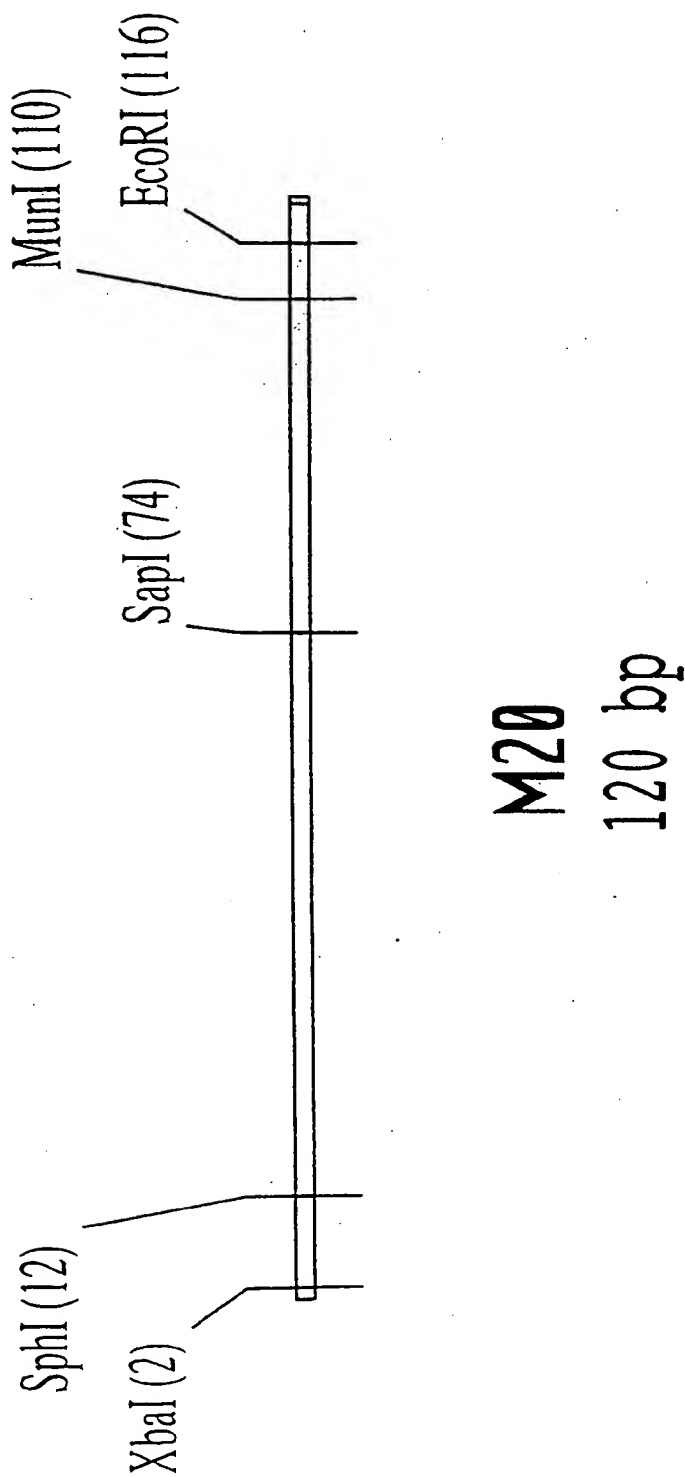


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 20:

	XbaI	SphI	
	-----	-----	
1	TCTAGAGCAT	GCGTAGGAGA	AAATAAATG AAACAAGCA CTATTGCACT
	AGATCTCGTA	CGCATCCTCT	TTTATTTTAC TTTGTTTCGT GATAACGTGA
		SapI	
		-----	
51	GGCACTCTTA	CCGTTGCTCT	TCACCCCTGT TACCAAAGCC GACTACAAAG
	CCGTGAGAAAT	GGCAACGAGA	AGTGGGACA ATGGTTTCGG CTGATGTTTC
	MunI	EcoRI	
	-----	-----	
101	ATGAAGTGCA	ATTGGAATTC	
	TACTTCACGT	TAACCTTAAG	

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

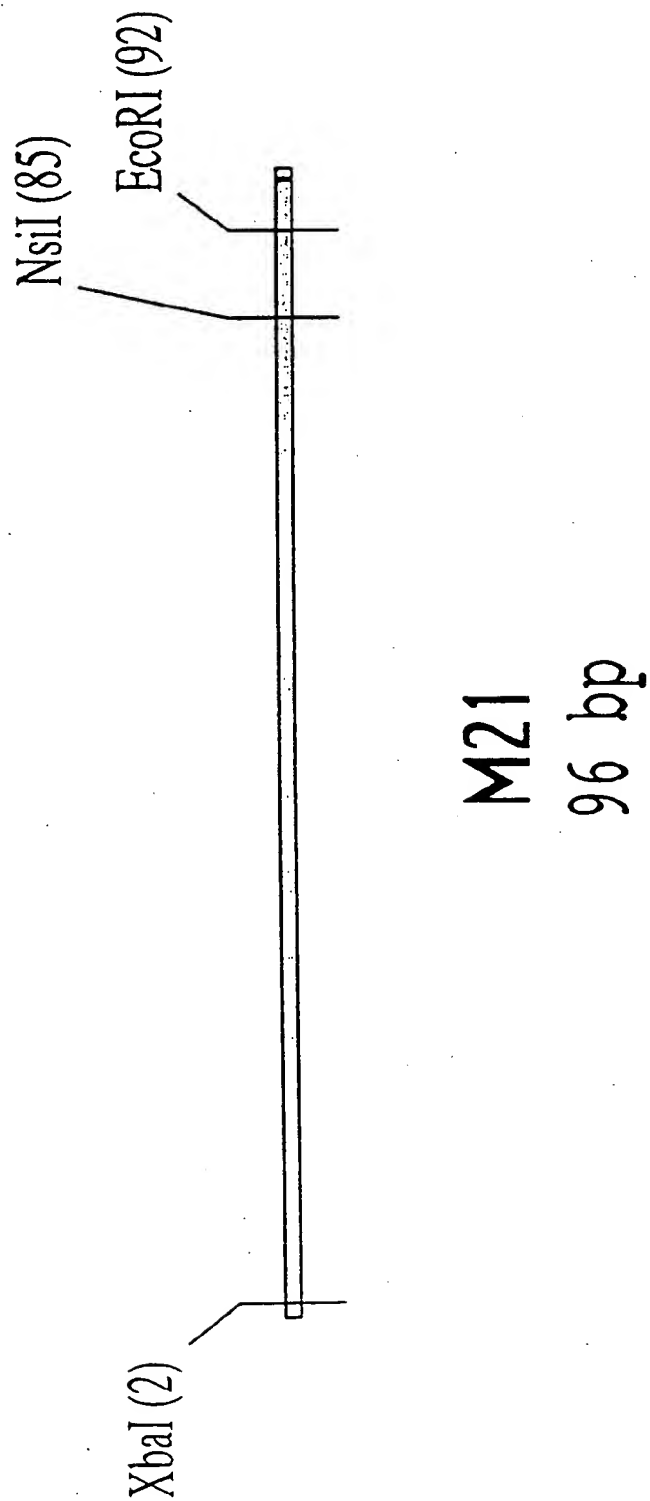


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 21:

XbaI

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1 TCTAGAGGTT GAGGTGATTT TATGAAAAG AATATCGCAT TTCTTCTTGC
 AGATCTCCAA CTCCTACTAA ATACTTTTTC TTATAGCGTA AAGAAGAACG

NsiI EcoRI

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51 ATCTATGTTT GTTTTCTTA TTGCTACAAA TGCATACGCT GAATTC  
 TAGATACAAG CAAAAAAGAT AACGATGTTT ACGTATGCCA CTTAAG

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

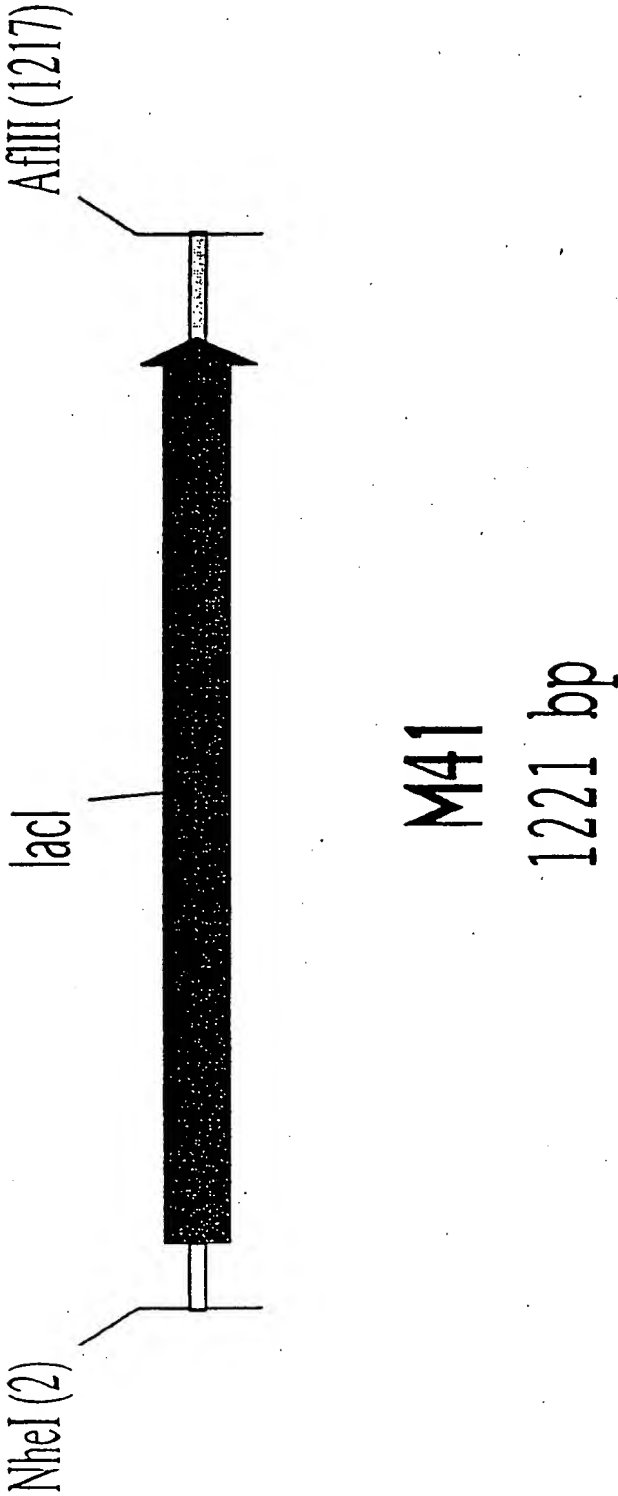


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 41:

NheI

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-----
1  GCTAGCATCG AATGGCGCAA AACCTTTCGC GGTATGGCAT GATAGCGCCC
   CGATCGTAGC TTACCGCGTT TTGGAAAGCG CCATACCGTA CTATCGCGGG

51  GGAAGAGAGT CAATTCAGGG TGGTGAATGT GAAACCAGTA ACGTTATACG
   CCTTCTCTCA GTTAAGTCCC ACCACTTACA CTTTGGTCAT TGCAATATGC

101 ATGTCGCAGA GTATGCCGGT GTCTCTTATC AGACCGTTTC CCGCGTGGTG
   TACAGCGTCT CATACGGCCA CAGAGAATAG TCTGGCAAAG GGCGCACCCAC

151 AACCAGGCCA GCCACGTTTC TGCGAAACG CGGGAACAAAG TGGAAGCGGC
   TTGGTCCGGT CGGTGCAAAG ACGCTTTTGC GCCCTTTTTC ACCTTCGCCC

201 GATGGCGGAG CTGAATTACA TTCCCTAACCG CGTGGCACAA CAACTGGCGG
   CTACCGCCCTC GACTTAATGT AAGGATTGGC GCACCGTGTT GTTGACCGCC

251 GCAAACAGTC GTTGCTGATT GGCGTTGCCA CCTCCAGTCT GGCCCTGCAC
   CGTTTGTGAG CAACGACTAA CCGCAACGGT GGAGGTCAGA CCGGACGTG

301 GCGCCGTCGC AAATTGTCGC GGCGATTAAA TCTCGCGCCG ATCAACTGGG
   CGCGGCAGCG TTTAACAGCG CCGCTAATTT AGAGCGCGGC TAGTTGACCC

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

351	TGCCAGCGTG	GTCGTGTCGA	TGGTAGAACG	AAGCGGCGTC	GAAGCCTGTA
	ACGGTCGCAC	CAGCACAGCT	ACCATCTTGC	TTGCGCGCAG	CTTCGGACAT
401	AAGCGGCGGT	GCACAAATCTT	CTCGCGCAAC	GTGTCAGTGG	GCTGATTATT
	TTCGCCGCCA	CGTGTTAGAA	GAGCGCGTTG	CACAGTCACC	CGACTAATAA
451	AACTATCCGC	TGGATGACCA	GGATGCTATT	GCTGTGGAAG	CTGCCTGCAC
	TTGATAGGCG	ACCTACTGGT	CCTACGATAA	CGACACCTTC	GACGGACGTG
501	TAATGTTCCG	GCGTTATTTC	TTGATGTCTC	TGACCAGACA	CCCATCAACA
	ATTACAAGGC	CGCAATAAAG	AACTACAGAG	ACTGGTCTGT	GGGTAGTTGT
551	GTATTATTTT	CTCCCATGAG	GACGGTACGC	GACTGGGCGT	GGAGCATCTG
	CATAATAAAA	GAGGGTACTC	CTGCCATGCG	CTGACCCGCA	CCTCGTAGAC
601	GTCGCCATTGG	GCCACCAGCA	AATCGCGCTG	TTAGCTGGCC	CATTAAAGTTC
	CAGCGTAACC	CGGTGGTTCGT	TTAGCGCGGAC	AATCGACCCG	GTAATTCAAG
651	TGTCTCGGCG	CGTCTGCGTC	TGGCTGGCTG	GCATAAATAT	CTCACTCGCA
	ACAGAGCCGC	GCAGACGCAG	ACCGACCGAC	CGTATTTATA	GAGTGAGCGT
701	ATCAAATTCA	GCCGATAGCG	GAACGGGAAG	GCGACTGGAG	TGCCATGTCC
	TAGTTTAAGT	CGGCTATCGC	CTTGCCCCCTC	CGCTGACCTC	ACGGTACAGG

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

751	GGTTTCAAC	AAACCATGCA	AATGCTGAAT	GAGGGCATCG	TTCCCCACTGC
	CCAAAAGTTG	TTTGGTACGT	TTACGACTTA	CTCCCCGTAGC	AAGGGTGACG
801	GATGCTGGTT	GCCAACGATC	AGATGGCGCT	GGGCGCAATG	CGTGCCATTA
	CTACGACCAA	CGGTTGCTAG	TCTACCGCGA	CCCGCGTTAC	GCACGGTAAT
851	CCGAGTCCGG	GCTGCGCGTT	GGTGCGGACA	TCTCGGTAGT	GGGATACGAC
	GGCTCAGGCC	CGACGCGCAA	CCACGCCCTGT	AGAGCCATCA	CCCTATGCTG
901	GATACCGAGG	ACAGCTCATG	TTATATCCCG	CCGCTGACCA	CCATCAAACA
	CTATGGCTCC	TGTCGAGTAC	AATATAGGC	GGCGACTGGT	GGTAGTTTGT
951	GGATTTTCGC	CTGCTGGGGC	AAACCAGCGT	GGACCGCTTG	CTGCAACTCT
	CCTAAAAGCG	GACGACCCCG	TTTGGTCGCA	CCTGGCGAAC	GACGTTGAGA
1001	CTCAGGGCCA	GGCGGTGAAG	GGCAATCAGC	TGTTGCCCCGT	CTCACTGGTG
	GAGTCCCGGT	CCGCCACTTC	CCGTTAGTCG	ACAACGGGCA	GAGTGACCAC
1051	AAAAGAAAAA	CCACCCTGGC	TCCCAATACG	CAAACCGCCT	CTCCCCGCGC
	TTTTCTTTTT	GGTGGGACCG	AGGGTTATGC	GTTTGGCGGA	GAGGGGCGCG
1101	GTTGGCCGAT	TCACTGATGC	AGCTGGCACG	ACAGGTTTCC	CGACTGGAAA
	CAACCGGCTA	AGTGACTACG	TCGACCGTGC	TGTCCAAAGG	GCTGACCCTT

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

1151 GCGGGCAGTG AGGCTACCCG ATAAAGCGG CTTCCCTGACA GGAGGCCCGTT  
CGCCCGTCAC TCCGATGGGC TATTTTCGCC GAAGGACTGT CCTCCGGCAA

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1201 TTGTTTGTCA GCCCACTTAA G
AACAAACGT CGGTGAATT C

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

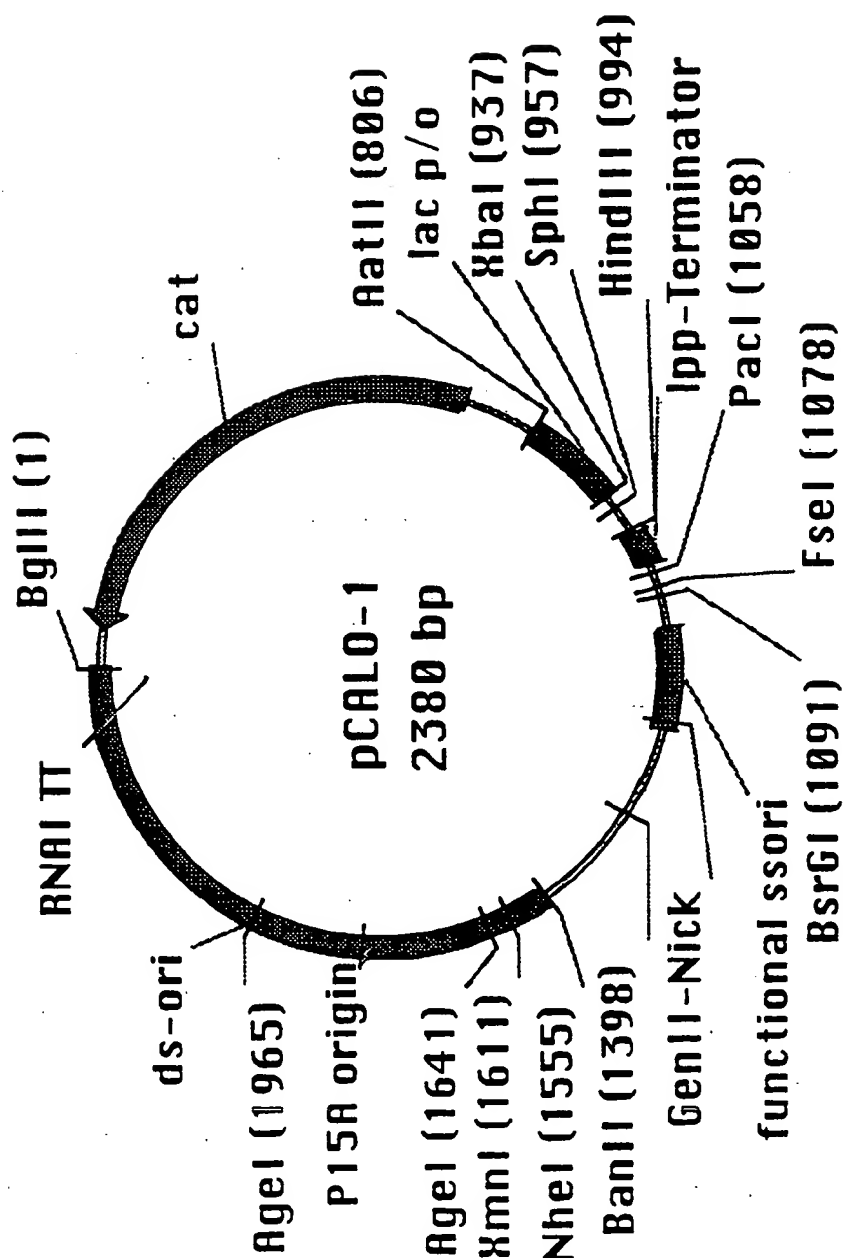


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

pCALO-1:
 BglII
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1	GATCTAGCAC	CAGCGGTTTA	AGGGCACCAA	TAACTGCCCTT	AAAAAATTA
	CTAGATCGTG	GTCCGCAAAAT	TCCCGTGGTT	ATTGACGGAA	TTTTTTTAAAT
51	CGCCCCGCC	TGCCACTCAT	CGCAGTACTG	TTGTAATTCA	TTAAGCATTC
	GCGGGCGGG	ACGGTGAGTA	GCGTCATGAC	AACATTAAGT	AATTCGTAAG
101	TGCCGACATG	GAAGCCATCA	CAAACGGCAT	GATGAACCTG	AATCGCCAGC
	ACGGCTGTAC	CTTCGGTAGT	GTTTGCCGTA	CTACTTGGAC	TTAGCGGTCG
151	GGCATCAGCA	CCTTGTCGCC	TTGCGTATAA	TATTTGCCCA	TAGTGAAAAC
	CCGTAGTCGT	GGAACAGCGG	AACGCATATT	ATAAACGGGT	ATCACTTTTG
201	GGGGGCCGAAG	AAGTTGTCCA	TATTGGCTAC	GTTTAAATCA	AAACTGGTGA
	CCCCCGCTTC	TTCAACAGGT	ATAACCGATG	CAAATTTAGT	TTTGACCACT
251	AACTCACCCA	GGGATTGGCT	GAGACGAAA	ACATATTCTC	AATAAACCCCT
	TTGAGTGGGT	CCCTAACCGA	CTCTGCTTTT	TGTATAAGAG	TTATTTTGGGA
301	TTAGGGAAAT	AGGCCAGGTT	TTCACCCGTAA	CACGCCACAT	CTTGCGAATA
	AATCCCCTTA	TCCGGTCCAA	AAGTGGCATT	GTGCGGTGTA	GAACGCTTAT

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

351	TATGTGTAGA AACTGCCGGA AATCGTCGTG GTATTCACCTC CAGAGCGATG ATACACATCT TTGACGGCCT TTAGCAGCAC CATAAGTGAG GTCTCGCTAC
401	AAAACGTTTC AGTTTGCTCA TGGAAAACGG TGTAACAAGG GTGAACACTA TTTTGCAAG TCAAACGAGT ACCTTTTGCC ACATTGTTCC CACTTGTGAT
451	TCCCATATCA CCAGCTCACC GTCCTTCATT GCCATACGGA ACTCCGGGTG AGGGTATAGT GGTCGAGTGG CAGAAAGTAA CGGTATGCCT TGAGGCCCCAC
501	AGCATTTCATC AGCGGGGCAA GAATGTGAAT AAAGGCCGGA TAAAACTTGT TCGTAAGTAG TCCGCCCGGT CTTACACTTA TTTCCGGCCT ATTTGAACA
551	GCTTATTTT CTTACGGTC TTTAAAAAGG CCGTAATATC CAGCTGAACG CGAATAAAAA GAAATGCCAG AAATTTTCC GCATTATAG GTCGACTTGC
601	GTCTGGTTAT AGGTACATTG AGCAACTGAC TGAAATGCCCT CAAAATGTTT CAGACCAATA TCCATGTAAC TCGTTGACTG ACTTTACGGA GTTTTACAAG
651	TTTACGATGC CATTGGGATA TATCAACGGT GGTATATCCA GTGATTTTTT AAATGCTACG GTAACCCCTAT ATAGTTGCCA CCATATAGGT CACTAAAAAA
701	TCTCCATTT AGCTTCCTTA GCTCCTGAAA ATCTCGATAA CTCAAAAAAT AGAGGTAAAA TCGAAGGAAT CGAGGACTTT TAGAGCTATT GAGTTTTTA

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

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751  ACGCCCGGTA GTGATCTTAT TTCATTATGG TGAAAGTTGG AACCTCACCC
    TCGGGGCCAT CACTAGAATA AAGTAATACC ACTTCAACC TTGGAGTGGG

    AatII
    ~~~~~
801 GACGTCTAAT GTGAGTTAGC TCACTCATTA GGCACCCAG GCTTTACACT
 CTGCAGATTA CACTCAATCG AGTGAGTAAT CCGTGGGTC CGAAATGTGA

851 TTATGCTTCC GGCTCGTATG TTGTGTGGAA TTGTGAGCGG ATAACAATTT
 AATACGAAGG CCGAGCATAC AACACACCTT AACACTCGCC TATTGTTAAA

 XbaI
                                ~~~~~
901  CACACAGGAA ACAGCTATGA CCATGATTAC GAATTCTAG ACCCCCCCCC
    GTGTGTCCTT TGTCGATACT GGTACTAATG CTTAAAGATC TGGGGGGGGG

                                SphI
                                ~~~~~
951 CGCATGCCAT AACTTCGTAT AATGTACGCT ATACGAAGTT ATAAGCTTGA
 GCGTACGGTA TTGAAGCATA TTACATGCCA TATGCTTCAA TATTCGAACT

1001 CCTGTGAAGT GAAAATGGC GCAGATTGTG CGACATTTT TTTGTCTGCC
 GGACACTTCA CTTTTTACCG CGTCTAACAC GCTGTAAAAA AAACAGACGG

 HindIII
                                ~~~~~

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

	PacI	FseI	BsrGI
	~~~~~	~~~~~	~~~~~
1051	GTTTAAATTAA AGGGGGGGG GGGCCGGCCT GGGGGGGGT GTACATGAAA CAAATTAATT TCCCCCCCCC CCGGCCGGA CCCCCCCCA CATGTACTTT		
1101	TTGTAAACGT TAAATTTTG TTAATAATTCG CGTTAAATTT TTGTAAATC AACATTTGCA ATTATAAAC AATTTTAAGC GCAATTTAAA AACAAATTTAG		
1151	AGCTCATTTT TTAACCAATA GGCCGAAATC GGCAAAATCC CTTATAAATC TCGAGTAAAA AATTGGTTAT CCGCTTTAG CCGTTTTAGG GAATATTTAG		
1201	AAAAGAATAG ACCGAGATAG GGTGAGTGT TGTTCCAGTT TGGAAACAAGA TTTCTCTATC TGGCTCTATC CCAACTCACA ACAAGGTCAA ACCTTGTTCT		
1251	GTCCACTATT AAAGAACGTG GACTCCAACG TCAAAGGGCG AAAAACCGTC CAGGTGATAA TTCTCTGCAC CTGAGGTTGC AGTTTCCCGC TTTTGGGCAG		
1301	TATCAGGGCG ATGGCCCACT ACGAGAACCA TCACCCCTAAT CAAGTTTTT ATAGTCCCCG TACCGGGTGA TGCTCTTGGT AGTGGGATTA GTTCAAAAAA		
			BanII
			~~~~~
1351	GGGGTCGAGG TGCCGTAAAG CACTAAATCG GAACCCCTAAA GGGAGCCCCC CCCCAGCTCC ACGGCATTTC GTGATTTAGC CTTGGGATTT CCCTCGGGGG		

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued).

	GATGCGAGCC	AGCAAGCTGA	CGCCGCTCGC	CTTTACCGAA	TGCTTGCCCC
1751	CGGAGATTTC	CTGGAAGATG	CCAGGAAGAT	ACTTAACAGG	GAAGTGAGAG
	GCCTCTAAAG	GACCTTCTAC	GGTCCCTTCTA	TGAATTGTCC	CTTCACTCTC
1801	GGCCGCGGCA	AAGCCGTTT	TCCATAGGCT	CCGCCCCCCT	GACAAGCATC
	CCGCGGCCGT	TTCGGCAAAA	AGGTATCCGA	GGCGGGGGA	CTGTTCTGTAG
1851	ACGAAATCTG	ACGCTCAAAT	CAGTGGTGGC	GAAACCCGAC	AGGACTATAA
	TGCTTTAGAC	TGCGAGTTTA	GTCACCACCG	CTTTGGGCTG	TCCTGATATT
1901	AGATACCAGG	CGTTTCCCCC	TGGCGGCTCC	CTCCTGCGCT	CTCCTGTTCC
	TCTATGGTCC	GCAAAGGGG	ACCGCCGAGG	GAGGACGCCA	GAGGACAAGG
	AgeI ~~~~~				
1951	TGCCTTTCCG	TTTACCAGTG	TCATTCCGCT	GTTATGGCCG	CGTTTGCTCTC
	ACGGAAAGCC	AAATGGCCAC	AGTAAGGCCA	CAATACCCGC	GCAAAACAGAG
2001	ATTCCACGCC	TGACACTCAG	TTCCGGGTAG	GCAGTTCGCT	CCAAGCTGGA
	TAAGGTGCGG	ACTGTAGTC	AAGCCCCATC	CGTCAAGCCA	GGTTCGACCT
2051	CTGTATGCAC	GAACCCCCCG	TTCAGTCCGA	CCGCTGCGCC	TTATCCGGTA
	GACATACGTG	CTTGGGGGGC	AAGTCAGGCT	GGCGACGCGG	AATAGGCCAT

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

1401	GATTAGAGC	TTGACGGGGA	AAGCCGGCGA	ACGTGGCGAG	AAAGGAAGGG
	CTAAATCTCG	AACTGCCCCCT	TTCGGCCCGCT	TGCACCGCTC	TTTCCCTTCCC
1451	AAGAAAGCGA	AAGGAGCGGG	CGCTAGGGCG	CTGGCAAGTG	TAGCGGTCAC
	TTCTTTTCGCT	TTCCCTCGCCC	GCGATCCCCG	GACCGTTCAC	ATCGCCAGTG
1501	GCTGCGCGTA	ACCACCACAC	CCGCCGCGCT	TAATGCGCCG	CTACAGGGCG
	CGACGCGCAT	TGGTGGTGTG	GGCGGCGCGA	ATTACGCGGC	GATGTCCCCG
NheI					
	~~~~~				
1551	CGTGCTAGCG	GAGTGTATAC	TGGCTTACTA	TGTTGGCACT	GATGAGGGTG
	GCACGATCGC	CTCACATATG	ACCGAATGAT	ACAACCGTGA	CTACTCCCCAC
XmnI					
	~~~~~				
1601	TCAGTGAAGT	GCTTCATGTG	GCAGGAGAAA	AAAGGCTGCA	CCGGTGCGTC
	AGTCACCTCA	CGAAGTACAC	CGTCCTCTTT	TTTCCGACGT	GGCCACGCAG
1651	AGCAGAATAT	GTGATACAGG	ATATATTCCG	CTTCCTCGCT	CACTGACTCG
	TCGTCTTATA	CACTATGTCC	TATATAAGGC	GAAGGAGCGA	GTGACTGAGC
1701	CTACGCTCGG	TCGTTGCACT	GCGGCGAGCG	GAAATGGCTT	ACGAACGGGG
				AgeI	~~~~~

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

2101	ACTATCGTCT	TGAGTCCAAC	CCGGAAGAC	ATGCAAAAGC	ACCACTGGCA
	TGATAGCAGA	ACTCAGGTG	GGCCTTCTG	TACGTTTTCG	TGGTGACCGT
2151	GCAGCCACTG	GTAATTGATT	TAGAGGAGTT	AGTCTTGAAG	TCATGCGCCG
	CGTCGGTGAC	CATTAACTAA	ATCTCCTCAA	TCAGAACTTC	AGTACGCGGC
2201	GTTAAGGCTA	AACTGAAAGG	ACAAGTTTTA	GTGACTGCGC	TCCTCCAAGC
	CAATTCCGAT	TTGACTTTCC	TGTTCAAAAT	CACTGACGCG	AGGAGGTTTCG
2251	CAGTTACCTC	GGTTCAAAGA	GTTGGTAGCT	CAGAGAACCT	ACGAAAACC
	GTCAATGGAG	CCAAGTTTCT	CAACCATCGA	GTCTCTTGGA	TGCTTTTGG
2301	GCCCTGCAAG	GCGGTTTTTT	CGTTTTCAGA	GCAAGAGATT	ACGCCAGAC
	CGGGACGTC	CGCCAAAAAA	GCAAAAGTCT	CGTTCTCTAA	TGCCGCTCTG
BglII					
2351	CAAACGATC	TCAAGAAGAT	CATCTTATTA		
	GTTTGTCTAG	AGTTCTTCTA	GTAGAATAAT		

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

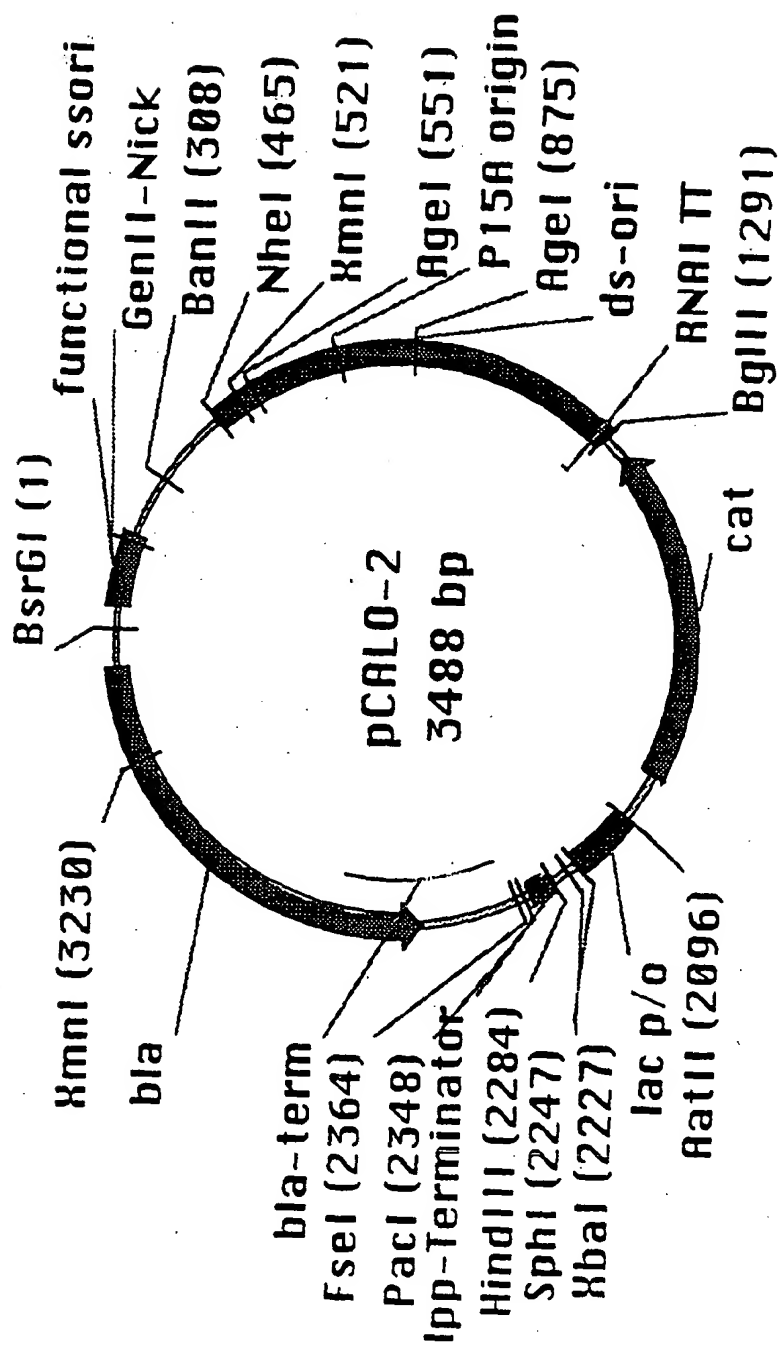




Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

## pCALO-2:

BsrGI

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| | | | | | |
|-----|-------------|-------------|------------|-------------|-------------|
| 1 | GTACATGAAA | TTGTAACGT | TAATATTTG | TTAAAAATTCG | CGTTAAATTT |
| | CATGTACTTT | AACATTGCA | ATTATAAAC | AATTTAAGC | GCAATTTAAA |
| 51 | TTGTTAATC | AGCTCATTTT | TTAACCAATA | GGCCGAAATC | GGCAAAATCC |
| | AACAATTTAG | TCGAGTAAAA | AATTGGTTAT | CCGGCTTTAG | CCGTTTTAGG |
| 101 | CTTATAAATC | AAAAGAAATAG | ACCGAGATAG | GGTTGAGTGT | TGTTCCAGTT |
| | GAATATTTAG | TTTTTCTTATC | TGGCTCTATC | CCAAC TCACA | ACAAGGTCAA |
| 151 | TGGAACAAGA | GTCCACTATT | AAAGAACGTG | GACTCCAACG | TCAAAGGGCG |
| | ACCTTGTTCT | CAGGTGATAA | TTTCTTGCAC | CTGAGGTTGC | AGTTTCCCCG |
| 201 | AAAACCCGTC | TATCAGGGCG | ATGGCCCACT | ACGAGAACCA | TCACCCCTAAT |
| | TTTTTTGGCAG | ATAGTCCCGC | TACCGGGTGA | TGCTCTTGGT | AGTGGGATTA |
| 251 | CAAGTTTTTT | GGGGTCGAGG | TGCCGTAAAG | CACTAAATCG | GAACCCTAAA |
| | GTTCAAAAAA | CCCCAGCTCC | ACGGCATTTT | GTGATTTAGC | CTTGGGATTT |
| 301 | GGGAGCCCCC | GATTAGAGC | TTGACGGGGA | AAGCCGGCGA | ACGTGGCGAG |

BanII

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

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CCCCTCGGGG  CTAAATCTCG  AACTGCCCCCT  TTCGGCCGCT  TGCACCGCTC

351  AAAGGAAGGG  AAGAAAGCGA  AAGAGCGGG  CGTAGGGCG  CTGGCAAGTG
      TTTCCTTCCC  TTCCTTCGCT  TTCCTCGCCC  GCGATCCCCG  GACCGTTCAC

401  TAGCGGTCAC  GTCGCGCGTA  ACCACCACAC  CCGCCGCGCT  TAATGCGCCG
      ATCGCCAGTG  CGACGCGCAT  TGGTGGTGTG  GCGGCGCGGA  ATTACGCGGC

      NheI
      ~~~~~
451 CTACAGGGCG CGTGCTAGCG GAGTGATAC TGGCTTACTA TGTGGCACT
 GATGTCCCGC GCACGATCGC CTCACATATG ACCGAATGAT ACAACCGTGA

 XmnI
      ~~~~~
      AgeI
501  GATGAGGGTG  TCAGTGAAGT  GCTTCATGTG  GCAGGAGAAA  AAAGGCTGCA
      CTACTCCCCAC  AGTCACTTCA  CGAAGTACAC  CGTCCTCTTT  TTTCCGACGT

      AgeI
      ~~~~~
551 CCGGTGCGTC AGCAGAATAT GTGATACAGG ATATATCCG CTTCCTCGCT
 GGCCACGCAG TCGTCTTATA CACTATGTCC TATATAAGGC GAAGGAGCGA

601 CACTGACTCG CTACGCTCGG TCGTTCGACT GCGGCGAGCG GAAATGGCTT

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

	GTGACTGAGC	GATGCGAGCC	AGCAAGCTGA	CGCCGCTCGC	CTTTACCGAA
651	ACGAACGGGG	CGGAGATTTC	CTGGAAGATG	CCAGGAAGAT	ACTTAACAGG
	TGCTTGCCCC	GCCTCTAAG	GACCTTCTAC	GTCCCTTCTA	TGAATTGTCC
701	GAAGTGAGAG	GGCCGCGGCA	AAGCCGTTT	TCCATAGGCT	CCGCCCCCCCT
	CTTCACTCTC	CCGGCGCCGT	TTCGGCAAAA	AGGTATCCGA	GGCGGGGGGA
751	GACAAGCATC	ACGAAATCTG	ACGCTCAAAT	CAGTGGTGGC	GAAACCCGAC
	CTGTTTCGTAG	TGCTTTAGAC	TGCGAGTTTA	GTCACCACCG	CTTTGGGCTG
801	AGGACTATAA	AGATACCAGG	CGTTTCCCCC	TGGCGGCTCC	CTCCTGCGCT
	TCCTGATATT	TCTATGGTCC	GCAAAGGGGG	ACCGCCGAGG	GAGACGCGA
			AgeI		
			~~~~~		
851	CTCCTGTTCC	TGCCTTTCGG	TTTACC GG TG	TCATTCCGCT	GTTATGGCCG
	GAGGACAAGG	ACGAAAGCC	AAATGGCCAC	AGTAAGGCGA	CAATACCGGC
901	CGTTTGTCTC	ATTCCACGCC	TGACACTCAG	TTCCGGGTAG	GCAGTTCGCT
	GCAAACAGAG	TAAGGTGCCG	ACTGTGAGTC	AAGGCCCATC	CGTCAAGCGA
951	CCAAGCTGGA	CTGTATGCAC	GAACCCCCCG	TTCAGTCCGA	CCGCTGCGCC
	GGTTCGACCT	GACATACGTG	CTTGGGGGGC	AAGTCAGGCT	GGCGACGCGG

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

1001	TTATCCGGTA	ACTATCGTCT	TGAGTCCAAC	CCGGAAGAC	ATGCAAAAGC
	AATAGGCCAT	TGATAGCAGA	ACTCAGGTG	GGCCTTCTG	TACGTTTTCG
1051	ACCACTGGCA	GCAGCCACTG	GTAATTGATT	TAGAGGAGTT	AGTCTTGAAG
	TGGTGACCGT	CGTCGGTGAC	CATTAACATA	ATCTCCTCAA	TCAGAACTTC
1101	TCATGCGCCG	GTTAAGGCTA	AACTGAAAGG	ACAAGTTTAA	GTGACTGCCG
	AGTACGCGGC	CAATTCCGAT	TTGACTTTCC	TGTTCAAAAT	CACTGACGCG
1151	TCCTCCAAGC	CAGTTACCTC	GGTTCAAAGA	GTTGGTAGCT	CAGAGAACCT
	AGGAGGTTCG	GTCAATGGAG	CCAAGTTTCT	CAACCATCGA	GTCTCTTGGA
1201	ACGAAAACC	GCCCTGCAAG	GCGGTTTTT	CGTTTTCAGA	GCAAGAGATT
	TGCTTTTGG	CGGACGTTT	CGCCAAAAA	GCAAAAGTCT	CGTTCTCTAA
BgIII					
1251	ACGCGCAGAC	CAAACGATC	TCAAGAAGAT	CATCTTATTA	GATCTAGCAC
	TGCGCGTCTG	GTTTTGCTAG	AGTTCTTCTA	GTAGAATAAT	CTAGATCGTG
1301	CAGCGGTTTA	AGGCAACCAA	TAACTGCCCT	AAAAAATA	CGCCCCGCC
	GTCCGCAAAT	TCCCGTGGTT	ATTGACGGAA	TTTTTTTAA	GCGGGCGGG

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

1351	TGCCACTCAT	CGCAGTACTG	TTGTAATTCA	TTAAGCATTC	TGCCGACATG
	ACGGTGAGTA	GCGTCATGAC	AACATTAAGT	AATTCGTAAG	ACGGCTGTAC
1401	GAAGCCATCA	CAAACGGCAT	GATGAACCTG	AATCGCCAGC	GGCATCAGCA
	CTTCGGTAGT	GTTTGCCGTA	CTACTTGAC	TTAGCGGTCG	CCGTAGTCGT
1451	CCTTGTCGCC	TTGCGTATAA	TATTTGCCCA	TAGTGAAAC	GGGGCGAAG
	GGAACAGCGG	AACGCATATT	ATAAACGGGT	ATCACTTTTG	CCCCCGCTTC
1501	AAGTTGTCCA	TATTGGCTAC	GTTTAAATCA	AAACTGGTGA	AACTCACCCA
	TTCAACACAGT	ATAACCGATG	CAAAATTAGT	TTTGACCACT	TTGAGTGGGT
1551	GGGATTGGCT	GAGACGAAA	ACATATTCTC	AATAAACCCCT	TTAGGGAAAT
	CCCTAACCGA	CTCTGCTTTT	TGTATAAGAG	TTATTTGGGA	AATCCCTTTA
1601	AGGCCAGGTT	TTCAACCGTAA	CACGCCACAT	CTTGCCGAATA	TATGTGTAGA
	TCCGGTCCAA	AAGTGGCATT	GTGCGGTGTA	GAACGCTTAT	ATACACATCT
1651	AACTGCCCGA	AATCGTCGTG	GTATTCACCTC	CAGAGCGATG	AAAACGTTC
	TTGACGGCCT	TTAGCAGCAC	CATAAGTGAG	GTCTCGCTAC	TTTTTGCAAAG
1701	AGTTTGCTCA	TGGAAAACGG	TGTAACAAGG	GTGAACACTA	TCCCATATCA
	TCAAACGAGT	ACCTTTTGCC	ACATTGTTCC	CACCTGTGAT	AGGGTATAGT

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

1751	CCAGCTCACC	GTCTTTCATT	GCCATACGGA	ACTCCGGGTG	AGCATTCATC
	GGTCGAGTGG	CAGAAAGTAA	CGGTATGCCT	TGAGGCCCCAC	TCGTAAAGTAG
1801	AGCGGGGCAA	GAATGTGAAT	AAAGGCCGGA	TAAAACTTGT	GCTTATTTTT
	TCCGCCCGTT	CTTACACTTA	TTTCCGGCCT	ATTTTGAACA	CGAATAAAAA
1851	CTTTACGGTC	TTTAAAAAGG	CCGTAATATC	CAGCTGAACG	GTCTGGTTAT
	GAAATGCCAG	AAATTTTCC	GGCATTATAG	GTCGACTTGC	CAGACCAATA
1901	AGGTACATTG	AGCAACTGAC	TGAAATGCCT	CAAAATGTTT	TTTACGATGC
	TCCATGTAA	TCGTTGACTG	ACTTTACGGA	GTTTACAAAG	AAATGCTACG
1951	CATTGGGATA	TATCAACGGT	GGTATATCCA	GTGATTTT	TCTCCATTTT
	GTAACCCCTAT	ATAGTTGCCA	CCATATAGGT	CACTAAAAAA	AGAGGTAAAA
2001	AGCTTCCTTA	GCTCCTGAAA	ATCTCGATAA	CTCAAAAAAT	ACGCCCGGTA
	TCGAAGGAAT	CGAGGACTTT	TAGAGCTATT	GAGTTTTTTA	TGCGGGCCAT
				AatII	
				~~~~~	
2051	GTGATCTTAT	TTCAATTATGG	TGAAAGTTGG	AACCTCACCC	GACGTCTAAT
	CACTAGAATA	AAGTAATACC	ACTTCAACC	TTGGAGTGGG	CTGCAGATTA
2101	GTGAGTTAGC	TCACTCATTA	GGCACCCCCAG	GCTTTACACT	TTATGCTTCC

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

	CACTCAATCG	AGTGAGTAAT	CCGTGGGGTC	CGAAATGTGA	AATACGAAGG
2151	GGCTCGTATG	TTGTGTGGAA	TTGTGAGCGG	ATAACAATTT	CACACAGGAA
	CCGAGCATAC	AACACACCTT	AACACTCGCC	TATTGTTAAA	GTGTGTCCTT
			XbaI		SphI
			~~~~~		~~~~~
2201	ACAGCTATGA	CCATGATTAC	GAATTTCTAG	ACCCCCCCC	CGCATGCCAT
	TGTCGATACT	GGTACTAATG	CTTAAAGATC	TGGGGGGGGG	GCGTACGGTA
				HindIII	
				~~~~~	
2251	AACTTCGTAT	AATGTACGCT	ATACGAAGTT	ATAAGCTTGA	CCTGTGAAGT
	TTGAAGCATA	TTACATGCGA	TATGCTTCAA	TATTCGAACT	GGACACTTCA
					PacI
					~~~~~
2301	GAAAAATGCG	GCAGATTGTG	CGACATTTT	TTTGTCTGCC	GTTTAATTAA
	CTTTTACC	CGTCTAACAC	GCTGTAAAAA	AAACAGACGG	CAAATTAAAT
				FseI	
			~~~~~		
2351	GGGGGGGGGC	CGGCCATTAT	CAAAAAGGAT	CTCAAGAAGA	TCCTTTGATC
	CCCCCCCCCG	GCCGGTAATA	GTTTTTCCCTA	GAGTTCTTCT	AGGAAACTAG

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

2401	TTTTTCTACGG	GGTCTGACGC	TCAGTGGAAC	GAAAACTCAC	GTAAAGGGAT
	AAAAGATGCC	CCAGACTGCG	AGTCACCTTG	CTTTTGAGTG	CAATTCCCTA
2451	TTTGGTCATG	AGATTATCAA	AAAGGATCTT	CACCTAGATC	CTTTTAAATT
	AAACCAGTAC	TCTAATAGTT	TTTCCTAGAA	GTGGATCTAG	GAAAAATTAA
2501	AAAATGAAG	TTTTAAATCA	ATCTAAAGTA	TATATGAGTA	AACTTGGTCT
	TTTTTACTTC	AAAATTTAGT	TAGATTTTCAT	ATATACTCAT	TTGAACCAGA
2551	GACAGTTACC	CAATGCTTAA	TCAGTGAGGC	ACCTATCTCA	GCGATCTGTC
	CTGTCAATGG	GTTACGGAATT	AGTCACTCCG	TGGATAGAGT	CGCTAGACAG
2601	TATTTTCGTC	ATCCATAGTT	GCCTGACTCC	CCGTCGTGTA	GATAACTACG
	ATAAAGCAAG	TAGGTATCAA	CGGACTGAGG	GGCAGCACAT	CTATTGATGC
2651	ATACGGGAGG	GCTTACCATC	TGGCCCCAGT	GCTGCAATGA	TACCGCGAGA
	TATGCCCTCC	CGAATGGTAG	ACCGGGGTCA	CGACGTTACT	ATGGCGCTCT
2701	CCCACGCTCA	CCGGCTCCAG	ATTATCAGC	AATAAACCAG	CCAGCCGGAA
	GGGTGCGAGT	GGCCGAGGTC	TAAATAGTCG	TTATTGGTC	GGTCGGCCCT
2751	GGGCCGAGCG	CAGAAAGTGGT	CCTGCAACTT	TATCCGCCCTC	CATCCAGTCT
	CCCGGCTCGC	GTCTTCACCA	GGACGTTGAA	ATAGCGGAG	GTAGGTCAGA



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

2801	ATTAACCTGTT GCCGGGAAGC TAGAGTAAGT AGTTCGCCAG TTAATAGTTT TAATTGACAA CGGCCCTTCG ATCTCATTTCA TCAAGCGGTC AATTATCAAA
2851	GCGCAACGTT GTTGCCATTG CTACAGGCAT CGTGGTGTCA CGCTCGTCGT CGCGTTGCAA CAACGGTAAC GATGTCCGTA GCACCACAGT GCGAGCAGCA
2901	TTGGTATGGC TTCATTTCAGC TCCGGTTCCC AACGATCAAG GCGAGTTACA AACCATAACG AAGTAAAGTCG AGGCCAAGGG TTGCTAGTTC CGCTCAATGT
2951	TGATCCCCCA TGTTGTGCAA AAAAGCGGTT AGCTCCCTTCG GTCCCTCCGAT ACTAGGGGGT ACAACACGTT TTTTCGCCCAA TCGAGGAAGC CAGGAGGCTA
3001	CGTTGTCAGA AGTAAAGTTGG CCGCAGTGTT ATCACTCATG GTTATGGCAG GCAACAGTCT TCATTCAACC GCGGTCACAA TAGTGAGTAC CAATACCGTC
3051	CACTGCATAA TTCTCTTACT GTCATGCCAT CCGTAAAGATG CTTTTCCTGTG GTGACGTATT AAGAGAAATGA CAGTACGGTA GGCAATCTAC GAAAAGACAC
3101	ACTGGTGAGT ACTCAACCAA GTCATTCTGA GAATAGTGTA TGCGGCGGACC TGACCACTCA TGAGTTGGTT CAGTAAAGACT CTTATCACAT ACGCCGCTGG
3151	GAGTTGCTCT TGCCCCGGCGT CAATACGGGA TAATACCGCG CCACATAGCA CTCAACGAGA ACGGGCCGCA GTTATGCCCT ATTATGGCGC GGTGTATCGT

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

XmnI	
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3201	GAAC TT TAAA AGTGCTCATC ATTGGAAAAC GTTCTTCGGG GCGAAAATC CTTGAAATTT TCACGAGTAG TAACCTTTTG CAAGAAGCCC CGCTTTTGAG
3251	TCAAGGATCT TACCGCTGTT GAGATCCAGT TCGATGTAAAC CCACTCGCGC AGTTCCCTAGA ATGGCGACAA CTCTAGGTCA AGCTACATTG GTGAGCGCG
3301	ACCCAACTGA TCCTCAGCAT CTTTACTTT CACCAGCGTT TCTGGGTGAG TGGGTTGACT AGGAGTCGTA GAAATGAAA GTGGTCGCAA AGACCCACTC
3351	CAAAACACAG AAGCAAAT GCCGCAAAA AGGGAATAAG GCGACACGG GTTTTTGTC TTCCGTTTTA CGCGTTTTT TCCCTTATTC CCGCTGTGCC
3401	AAATGTTGAA TACTCATACT CTCCTTTT CAATATTATT GAAGCATTTA TTTACAACCTT ATGAGTATGA GAAGGAAAA GTTATAATAA CTTCGTAAAT
BsrGI	
3451	TCAGGGTTAT TGTCTCATGA GCGGATACAT ATTTGAAT AGTCCCAATA ACAGAGTACT CGCCTATGTA TAAACTTA

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

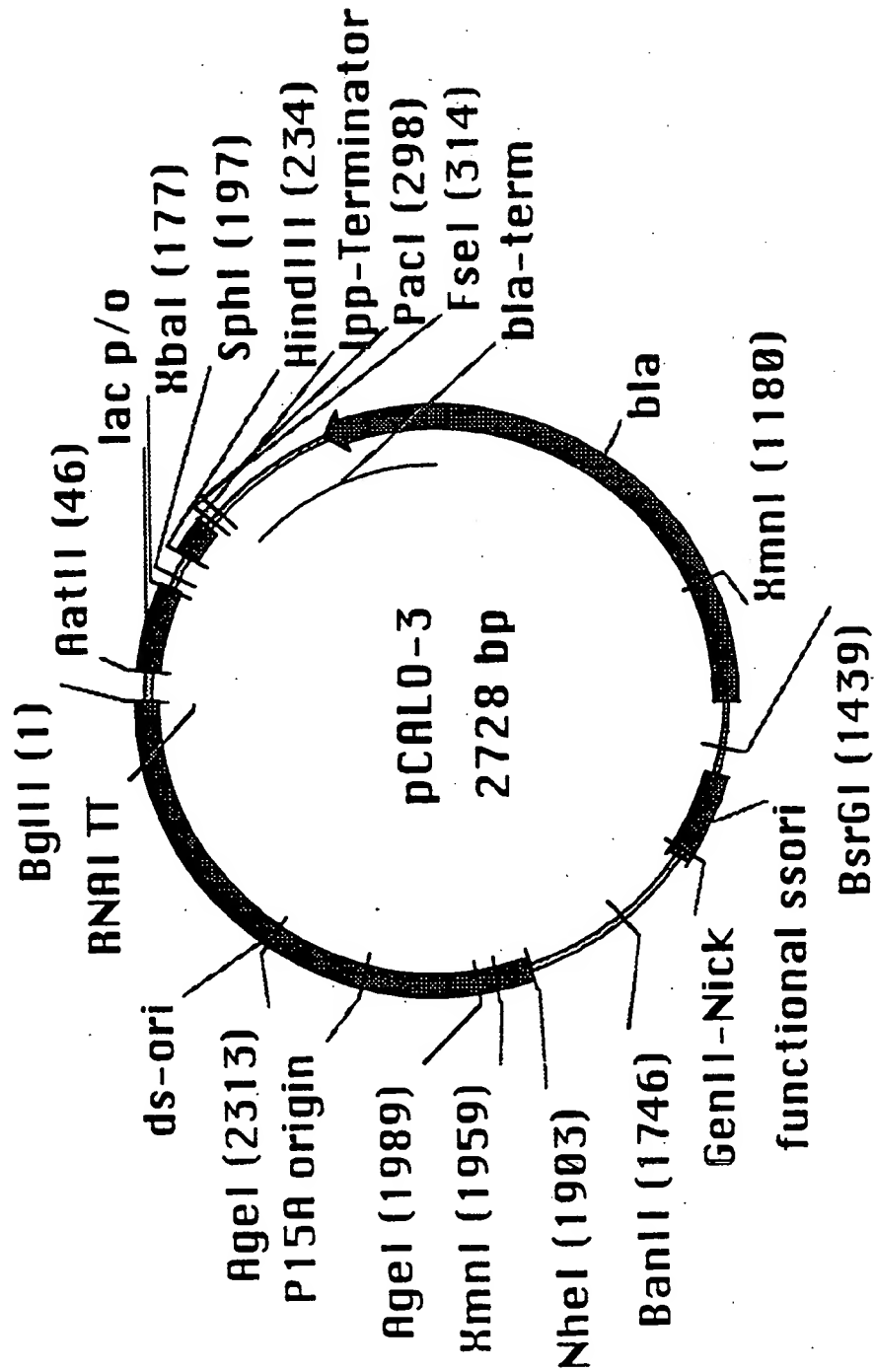


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

pCALO-3:		AatII	
		BglII	~~~~~
1	GATCTCATAA CTTCGTATAA TGTATGCTAT ACGAAGTTAT GACGTCTAAT		
	CTAGAGTATT GAAGCATATT ACATACGATA TGCTTCAATA CTGCAGATTA		
51	GTGAGTTAGC TCACTCATTA GGCACCCCAG GCTTTACACT TTATGCTTCC		
	CACTCAATCG AGTGAGTAAT CCGTGGGGTC CGAAATGTGA AATACGAAGG		
101	GGCTCGTATG TTGTGTGGAA TTGTGAGCGG ATAACAATTT CACACAGGAA		
	CCGAGCATAC AACACACCTT AACACTCGCC TATTGTTAA GTGTGTCCTT		
		XbaI	SphI
		~~~~~	~~~~~
151	ACAGCTATGA CCATGATTAC GAATTCTAG ACCCCCCCCC CGCATGCCAT		
	TGTCGATACT GGTACTAATG CTTAAAGATC TGGGGGGGGG GCGTACGGTA		
		HindIII	
		~~~~~	
201	AAC TTCGTAT AATGTACGCT ATACGAAGTT ATAAGCTTGA CCTGTGAAGT		
	TTGAAGCATA TTACATGCCA TATGCTTCAA TATTCGAACT GGACACTTCA		
			PacI

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

251	GAAAAATGGC GCAGATTGTG CGACATTTT TTTGTCTGCC GTTTAATTAA CTTTTACC CGTCTAACAC GCTGTAAAAA AACACAGACGG CAAATTAAAT	~~~~~
FseI		
301	GGGGGGGGC CGCCATTAT CAAAAGGAT CTCAAGAAGA TCCTTTGATC CCCCCCCCG GCGGTAATA GTTTTCCCTA GAGTTCTTCT AGGAAACTAG	~~~~~
351	TTTTCTACGG GGTCTGACGC TCAGTGGAAC GAAAACCTCAC GTTAAGGGAT AAAAGATGCC CCAGACTGCG AGTCACCTTG CTTTGTAGTG CAATTCCCCTA	
401	TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTAAATT AAACCAGTAC TCTAATAGTT TTTCCCTAGAA GTGGATCTAG GAAAAATTAA	
451	AAAAATGAAG TTTTAAATCA ATCTAAAGTA TATATGAGTA AACTTGGTCT TTTTTACTTC AAAATTTAGT TAGATTTCAT ATATACTCAT TTGAACCCAGA	
501	GACAGTTACC CAATGCTTAA TCAGTGAGGC ACCTATCTCA GCGATCTGTC CTGTCAATGG GTTACGAAAT AGTCACTCCG TGGATAGAGT CGCTAGACAG	
551	TATTTCGTTT ATCCATAGTT GCCTGACTCC CCGTCGTGTA GATAACTACG ATAAAGCAAG TAGGTATCAA CGGACTGAGG GGCAGCACAT CTATTGATGC	

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

601	ATACGGGAGG	GCTTACCATC	TGGCCCCCAGT	GCTGCAATGA	TACCGCGAGA
	TATGCCCTCC	CGAATGGTAG	ACCGGGGTCA	CGACGTTACT	ATGGCGCTCT
651	CCCACGCTCA	CCGGCTCCAG	ATTTATCAGC	AATAAACCCAG	CCAGCCGGAA
	GGTGCGAGT	GGCCGAGGTC	TAAATAGTCG	TTATTTGGTC	GGTCGGCCCTT
701	GGGCCGAGCG	CAGAGTGGT	CCTGCAACTT	TATCCGCCCTC	CATCCAGTCT
	CCCGGCTCGC	GTCTTCACCA	GGACGTTGAA	ATAGCGCGAG	GTAGGTCAGA
751	ATTAACTGTT	GCCGGGAAGC	TAGAGTAAGT	AGTTCGCCCAG	TTAATAGTTT
	TAATTGACAA	CGGCCCTTCG	ATCTCATTCA	TCAAGCGGTC	AATTATCAAA
801	GCGCAACGTT	GTTGCCATTG	CTACAGGCAT	CGTGGTGTC	CGCTCGTCGT
	CGCGTTGCAA	CAACGGTAAC	GATGTCCGTA	GCACCACAGT	GCGAGCAGCA
851	TTGGTATGGC	TTCATTTCAGC	TCCGGTTCCC	AACGATCAAG	GCGAGTTACA
	AACCATAACCG	AAGTAAGTCG	AGGCCAAGGG	TTGCTAGTTC	CGCTCAATGT
901	TGATCCCCCA	TGTTGTGCAA	AAAAGCGGTT	AGTCCCTTCG	GTCCCTCCGAT
	ACTAGGGGGT	ACAACACGTT	TTTTTCGCCAA	TCGAGGAAGC	CAGGAGGCTA
951	CGTTGTCAGA	AGTAAGTTGG	CCGCAGTGTT	ATCACTCATG	GTTATGGCAG
	GCAACAGTCT	TCATTCAACC	GGCGTCACAA	TAGTGAGTAC	CAATACCGTC

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

1001	CACTGCATAA TTCTCTTACT GTCATGCCAT CCGTAAGATG CTTTCTGTG GTGACGTATT AAGAGAAATGA CAGTACGGTA GGCATTCTAC GAAAAGACAC
1051	ACTGGTGAGT ACTCAACCAA GTCATTTCTGA GAATAGTGTA TCGGGCGACC TGACCACTCA TGAGTTGGTT CAGTAAGACT CTTATCACAT ACGCCGCTGG
1101	GAGTTGCTCT TGCCCGGCGT CAATACGGGA TAATACCGCG CCACATAGCA CTCAACGAGA ACGGCCGCA GTTATGCCCT ATTATGGCGC GGTGTATCGT
XmnI ~~~~~	
1151	GAACTTTAAA AGTGCTCATC ATTGGAAAAC GTTCTTCGGG GCGAAAAC TC CTTGAAAATT TCACGAGTAG TAACCTTTTG CAAGAAGCCC CGCTTTTGAG
1201	TCAAGGATCT TACCGCTGTT GAGATCCAGT TCGATGTAAC CCACTCGCGC AGTTCCCTAGA ATGGCGACAA CTCTAGGTCA AGCTACATTG GGTGAGCGCG
1251	ACCCAAC TGA TCCCTCAGCAT CTTTACTTT CACCAGCGTT TCTGGGTGAG TGGGTTGACT AGGAGTCGTA GAAAATGAAA GTGGTCGCAA AGACCCACTC
1301	CAAAAACAGG AAGGCAAAAT GCCGCAAAA AGGGAATAAG GCGACACGG GTTTTTGTCC TTCCGTTTTA CGCGTTTTT TCCCTTATTC CCGCTGTGCC
1351	AAATGTTGAA TACTCATACT CTTCCTTTTT CAATATTATT GAAGCATTTA

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

	TTTACAACCTT	ATGAGTATGA	GAAGGAAAAA	GTTATAATAA	CTTCGTAAAT
	BsrGI				
	~~~~~				
1401	TCAGGGTTAT	TGTCTCATGA	CGGATACAT	ATTTGAATGT	ACATGAAATT
	AGTCCCAATA	ACAGAGTACT	CGCCTATGTA	TAAACTTACA	TGTACTTTAA
1451	GTAACGTTA	ATATTTTGT	AAAATTCGCG	TTAAATTTT	GTTAAATCAG
	CATTGCAAT	TATAAAACAA	TTTTAAGCGC	AATTTAAAAA	CAATTTAGTC
1501	CTCATTTTTT	AACCAATAGG	CCGAAATCGG	CAAAATCCCT	TATAAATCAA
	GAGTAAAAAA	TTGGTTATCC	GGCTTTAGCC	GTTTTAGGGA	ATATTTAGTT
1551	AAGAATAGAC	CGAGATAGGG	TTGAGTGTTG	TTCCAGTTTG	GAACAAGAGT
	TTCTTATCTG	GCTCTATCCC	AACTCACAAAC	AAGGTCAAAC	CTTGTTCTCA
1601	CCACTATTAA	AGAACGTGGA	CTCCAACGTC	AAAGGGCGAA	AAACCGTCTA
	GGTGATAATT	TCTTGACACCT	GAGGTTGCAG	TTTCCCGCCTT	TTTGGCAGAT
1651	TCAGGGCGAT	GGCCCACTAC	GAGAACCATC	ACCCTAATCA	AGTTTTTGG
	AGTCCCGCTA	CCGGGTGATG	CTCTTGGTAG	TGGGATTAGT	TCAAAAAAAC
	BanII				
	~~~~~				



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

1701	GGTCGAGGTG CCGTAAAGCA CTAAATCGGA ACCCTAAAGG GAGCCCCCGA CCAGCTCCAC GGCATTTCGT GATTAGCCT TGGGATTTC CTCGGGGGCT	
1751	TTTAGAGCTT GACGGGGAAG GCCGGCGAAC GTGGCGAGAA AGGAAGGAA AAATCTCGAA CTGCCCCCTT CGCGCGCTTG CACCGCTCTT TCCTTCCCTT	
1801	GAAAGCGAAA GGAGCGGGCG CTAGGGCGCT GGCAAGTGTA GCGTCAACG CTTTCGCTTT CCTCGCCCCG GATCCCGCGA CCGTTCACAT CGCCAGTGCG	
1851	TGCGCGTAAC CACCACACCC GCCGCGCTTA ATGCGCCGCT ACAGGGCGCG ACGCGCATTG GTGGTGTGG CGCGCGGAAT TACGCGGCGA TGTCCCGCGC	
1901	TGCTAGCGGA GTGTATACTG GCTTACTATG TTGGCACTGA TGAGGGTGTC ACGATCGCCT CACATATGAC CGAATGATAC AACCGTGACT ACTCCCACAG	
1951	AGTGAAGTGC TTCATGTGGC AGGAGAAAAA AGGCTGCACC GGTGCGTCAG TCACTTCACG AAGTACACCG TCCTCTTTT TCCGACGTGG CCACGCAGTC	
2001	CAGAATATGT GATACAGGAT ATATTCCGCT TCCTCGCTCA CTGACTCGCT GTCTTATACA CTATGTCCTA TATAAGCGA AGGAGCGAGT GACTGAGCGA	

AgeI

XmnI

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

2051	ACGCTCGGTC GTTCGACTGC GCGAGCGGA AATGGCTTAC GAACGGGGCG TGCAGGCCAG CAAGCTGACG CCGCTCGCCT TTACCCGAATG CTTGCCCCCGC
2101	GAGATTTCCT GGAAGATGCC AGGAAGATAC TTAACAGGGA AGTGAGAGGG CTCTAAAGGA CCTTCTACGG TCCTTCTATG AATTGTCCCT TCACTCTCCC
2151	CCGCGGCAAA GCCGTTTTC CATAAGCTCC GCCCCCCTGA CAAGCATCAC GGCGCCGTTT CGGCAAAAAG GTATCCGAGG CGGGGGACT GTTCGTAGTG
2201	GAAATCTGAC GCTCAAATCA GTGGTGGCGA AACCCGACAG GACTATAAAG CTTTAGACTG CGAGTTTAGT CACCACCGCT TTGGGCTGTC CTGATATTTC
2251	ATACCAGGCG TTTCCCCCTG GCGGCTCCCT CCTGCGCTCT CCTGTTCCCTG TATGGTCCGC AAAGGGGGAC CGCCGAGGGA GGACGCGAGA GGACAAGGAC
AgeI ~~~~~	
2301	CCTTTCGGTT TACCGGTGTC ATCCGCTGT TATGGCCGCG TTTGTCTCAT GGAAAGCCAA ATGCCACAG TAAGCGACA ATACCGGCGC AACACAGTA
2351	TCCACGCCCTG AACTCAGTT CCGGGTAGGC AGTTCGCTCC AAGCTGGACT AGGTGCGGAC TGTGAGTCAA GGCCCATCCG TCAAGCGAGG TTCGACCTGA

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

2401	GTATGCACGA	ACCCCCCGTT	CAGTCCGACC	GCTGCGCCTT	ATCCGGTAAC
	CATACGTGCT	TGGGGGGCAA	GTCAGGCTGG	CGACGCGGAA	TAGGCCATTC
2451	TATCGTCTTG	AGTCCAACCC	GGAAAGACAT	GCAAAAGCAC	CACTGGCAGC
	ATAGCAGAAC	TCAGGTTGGG	CCTTCTCTGA	CGTTTTCGTG	GTGACCGTCG
2501	AGCCACTGGT	AATTGATTTA	GAGGAGTTAG	TCTTGAAGTC	ATGCGCCGGT
	TCGGTGACCA	TTAACTAAAT	CTCCTCAATC	AGAACTTCAG	TACGCGGCCA
2551	TAAGGCTAAA	CTGAAAGGAC	AAGTTTTAGT	GACTGCGCTC	CTCCAAGCCA
	ATTCCGATT	GACTTTCCTG	TTCAAAATCA	CTGACGCGAG	GAGGTTCCGT
2601	GTTACCTCGG	TTCAAAGAGT	TGGTAGCTCA	GAGAACCTAC	GAAAAACCGC
	CAATGGAGCC	AAGTTTCTCA	ACCATCGAGT	CTCTTGGATG	CTTTTGGCG
2651	CCTGCAAGGC	GGTTTTCG	TTTTTCAGAGC	AAGAGATTAC	GCGCAGACCA
	GGACGTTCCG	CCAAAAAAGC	AAAAGTCTCG	TTCTCTAATG	CGCGTCTGGT
Bg1II					
2701	AAACGATCTC	AAGAAGATCA	TCTTATTA		
	TTTGCTAGAG	TTCTTCTAGT	AGAATAAT		

Figure 35b: List of oligonucleotides used for synthesis of modules

M1: PCR using template

NoVspAatII: TAGACGTC

M2: synthesis

BloxA-A: TATGAGATCTCATAACTTCGTATAATGTACGCTATACG-  
AAGTTAT

BloxA-B: TAATAACTTCGTATAGCATAACATTATACGAAGTTATG-  
AGATCTCA

M3: PCR, NoVspAatII as second oligo

XloxS-muta: CATTTTTGCCCTCGTTATCTACGCATGCGATAACTTCGTA-  
TAGCGTACATTATACGAAGTTATTCTAGACATGGTCATAGCTGTTTCCTG

M7-I: PCR

gIII NEW-fow: GGGGGGAATTCGGTGGTGGTGGATCTGCGTGCGCTG-  
AAACGGTTGAAAGTTG

gIII NEW-rev: CCCCCCAAGCTTATCAAGACTCCTTATTACG

M7-II: PCR

gIII ss-fow: GGGGGGGGAATTCGGAGGCGGTTCCGGTGGTGGC

M7-III: PCR

gIII supernew-fow: GGGGGGGGAATTCGAGCAGAAGCTGATCTCT-  
GAGGAGGATCTGTAGGGTGGTGGCTCTGGTTCCGGTGATTTG

Figure 35b: List of oligonucleotides used for synthesis of modules (continued)

**M8: synthesis**

lox514-A: CCATAACTTCGTATAATGTACGCTATACGAAGTTATA

lox514-B: AGCTTATAACTTCGTATAGCGTACATTATACGAAGT-  
TATGGCATG

**M9II: synthesis**

M9II-fow: AGCTTGACCTGTGAAGTGAAAAATGGCGCAGATT-  
GTGCGACATTTTTTTTGTCTGCCGTTTAATTAAAGGGGGGGT

M9II-rev: GTACACCCCCCCCCAGGCCGGCCCCCCCCCCCCCTTTAA-  
TTAAACGGCAGACAAAAAAAATGTCGCACAATCTGCG

**M10II: assembly PCR with template**

bla-fow: GGGGGGGTGTACATTCAAATATGTATCCGCTCATG

bla-seq4: GGGTTACATCGAACTGGATCTC

bla1-muta: CCAGTTCGATGTAACCCACTCGCGCACCCAACTGATC-  
CTCAGCATCTTTACTTTCACC

blaII-muta: ACTCTAGCTTCCCGGCAACAGTTAATAGACTGGATG-  
GAGGCGG

bla-NEW: CTGTTGCCGGGAAGCTAGAGTAAG

bla-rev: CCCCCCTTAATTAAGGGGGGGGGCCGGCCATTATCAAA-  
AAGGATCTCAAGAAGATCC

**M11II/III: PCR, site-directed mutagenesis**

Figure 35b: List of oligonucleotides used for synthesis of modules (continued)

f1-fow: GGGGGGGGCTAGCACGCGCCCTGTAGCGGCGCATTA

f1-rev: CCCCCCTGTACATGAAATTGTAAACGTTAATATTTG

f1-t133.muta: GGGCGATGGCCCACTACGAGAACCATCACCTAATC

### M12: assembly PCR using template

p15-fow: GGGGGGAGATCTAATAAGATGATCTTCTTGAG

p15-NEWI: GAGTTGGTAGCTCAGAGAACCTACGAAAAACCGCCCTG-  
CAAGGCG

p15-NEWII: GTAGGTTCTCTGAGCTACCAACTC

p15-NEWIII: GTTCCCCCTGGCGGCTCCCTCCTGCGCTCTCCTGTTCT-  
GCC

p15-NEWIV: AGGAGGGAGCCGCCAGGGGGGAAAC

p15-rev: GACATCAGCGCTAGCGGAGTGTATAC

### M13: synthesis

BloxXB-A: GATCTCATAACTTCGTATAATGTATGCTATACGAAGTTA-  
TTCA

BloxXB-B: GATCTGAATAACTTCGTATAGCATACATTATACGAAGTTA-  
TGAGA

### M14-Ext2: PCR, site-directed mutagenesis

ColEXT2-fow: GGGGGGGGAGATCTGACCAAATCCCTTAACGTGAG

Col-mutal: GGTATCTGCGCTCTGCTGTAGCCAGTTACCTTCGG

Figure 35b: List of oligonucleotides used for synthesis of modules (continued)

Col-rev: CCCCCCGCTAGCCATGTGAGCAAAAGGCCAGCAA

M17: assembly PCR using template

CAT-1: GGGACGTCGGGTGAGGTCCAAC

CAT-2: CCATACGGAACCTCCGGGTGAGCATTATC

CAT-3: CCGGAGTTCCGTATGG

CAT-4: ACGTTTAAATCAAACTGG

CAT-5: CCAGTTTTGATTTAAACGTAGCCAATATGGACAACCTCTTC-  
GCCCCCGTTTTCACTATGGGCAAATATT

CAT-6: GGAAGATCTAGCACCAGGCGTTTAAG

M41: assembly PCR using template

LAC1: GAGGCCGGCCATCGAATGGCGCAAAAC

LAC2: CGCGTACCGTCCTCATGGGAGAAAATAATAC

LAC3: CCATGAGGACGGTACGCGACTGGGCGTGGAGCATCTGGTCGCA-  
TTGGGTCACCAGCAAATCCGCTGTTAGCTGGCCCATTAAG

LAC4: GTCAGCGGCGGGATATAACATGAGCTGTCCTCGGTATCGTCG

LAC5: GTTATATCCCGCCGCTGACCACCATCAAAC

LAC6: CATCAGTGAATCGGCCAACGCGCGGGGAGAGGCGGTTTGCGT4TTG-  
GGAGCCAGGGTGGTTTTTC

LAC7: GGTTAATTAACCTCACTGCCCCGCTTTCCAGTCGGGAAACCTGTCGTGCC-  
AGCTGCATCAGTGAATCGGCCAAC

M41-MCS-fow: CTAGACTAGTGTTTAAACCGGACCGGGGGGGGGGCTT-  
AAGGGGGGGGGGGGGG

Figure 35b: List of oligonucleotides used for synthesis of modules (continued)

M41-MCS-rev: CTAGCCCCCCCCCCTTAAGCCCCCCCCCGGTCCGGT-  
TTAAACACTAGT

M41-fow: CTAGACTAGTGTTTAAACGGACCGGGGGGGGGCTTAA-  
GGGGGGGGGGGGG

M41-rev: CCCCCCTTAAGTGGGCTGCAAAACAAAACGGCCTCC-  
TGTCAGGAAGCCGCTTTTATCGGGTAGCCTCACTGCCCGCTTCC

M41-A2: GTTGTTGTGCCACGCGGTTAGGAATGTAATTCAGCTCCGC

M41-B1: AACCGCGTGGCACAACAAC

M41-B2: CTCGTTCTACCATCGACACGACCACGCTGGCACCCAGTTG

M41-C1: GTGTCGATGGTAGAACGAAG

M41-CII: CCACAGCAATAGCATCCTGGTCATCCAGCGGATAGTT-  
AATAATCAGCCCCTGACACGTTGCGCGAG

M41-DI: GACCAGGATGCTATTGCTGTGG

M41-DII: CAGCGCGATTTGCTGGTGGCCCAATGCGACCAGATGC

M41-EI: CACCAGCAAATCGCGCTG

M41-EII: CCCGGACTCGGTAATGGCACGCATTGCGCCCAGCGCC

M41-FI: GCCATTACCGAGTCCGGG

#### M42: synthesis

Eco-H5-Hind-fow: AATTCCACCATCATCACCATTGACGTCTA

Eco-H5-Hind-rev: AGCTTAGACGTCAATGGTGATGATGGTGG



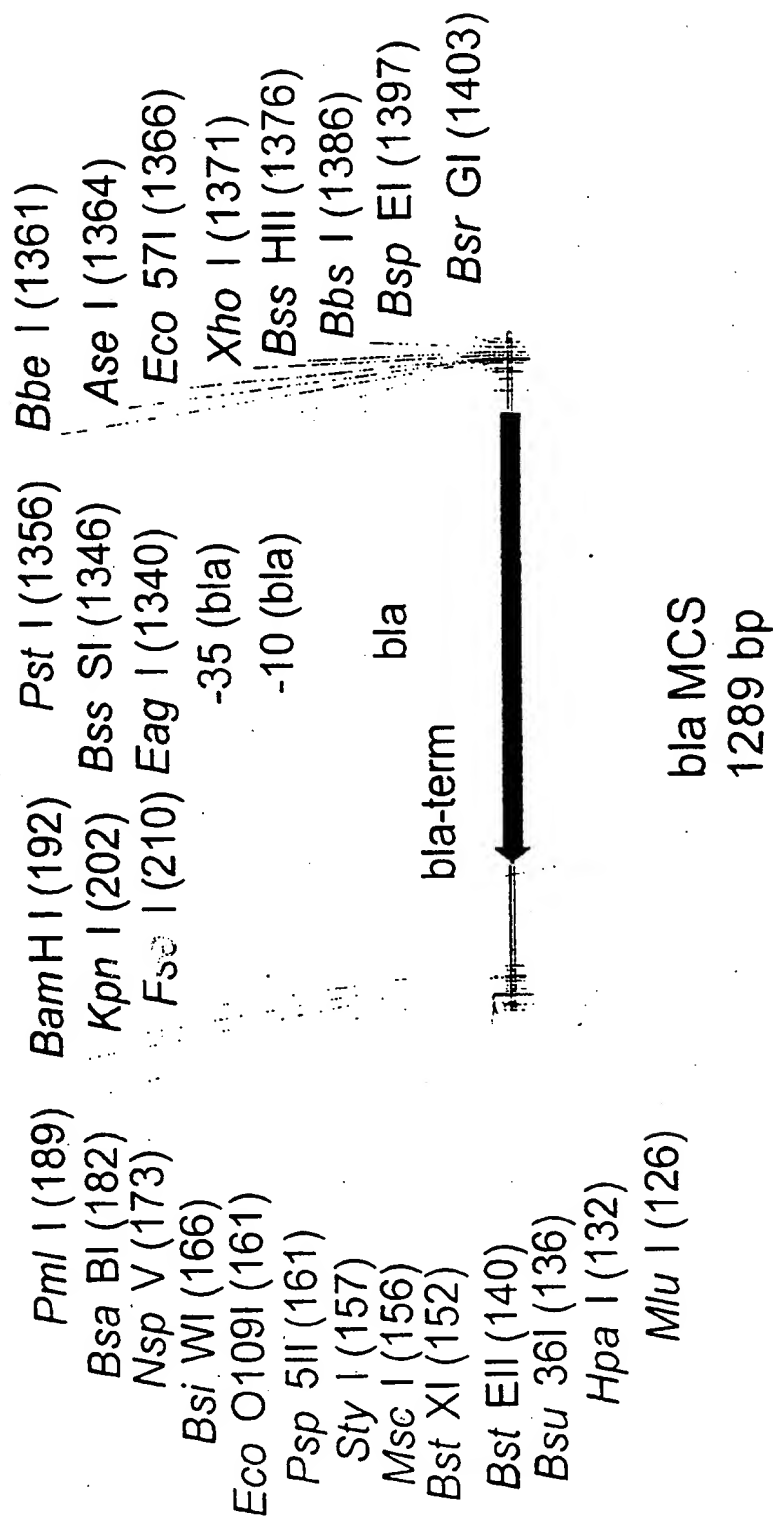
Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

		StyI			
		~~~~~			
		Psp5II			
		~~~~~			
		EcoO109I			
		~~~~~			
		BstXI			
		~~~~~			
		MscI			
		~~~~~			
		BsiWI NspV			
		~~~~~			
		MluI			
		~~~~~			
		Bsu36I			
		~~~~~			
		HpaI			
		~~~~~			
		BstEII			
		~~~~~			
		MscI			
		~~~~~			
126	CGCGTTAACC	TCAGGTGACC	AAGCCCCTGG	CCAAGTCCC	GTACGTTCCA
	CGGCAATTGG	AGTCCACTGG	TTCGGGACC	GGTCCAGGG	CATGCAAGCT
		PmlI			
		~~~~~			
		NspVBSaBI			
		~~~~~			
		BamHI			
		~~~~~			
		KpnI			
		~~~~~			
176	AGATTACCAT	CACGTGGATC	CGGTACCAGG	CCGGCCATTA	TCAAAAAGGA
	TCTAATGGTA	GTGCACCTAG	GCCATGGTCC	GGCCGGTAAT	AGTTTTTCCT
226	TCTCAAGAAG	ATCCTTTTGAT	CTTTTCTACG	GGGTCTGACG	CTCAGTGGAA
	AGAGTTCCTC	TAGGAAACTA	GAAAAGATGC	CCCAGACTGC	GAGTCACCTT
276	CGAAAACCTCA	CGTTAAGGGA	TTTTGGTCAT	GAGATTATCA	AAAAGGATCT
	GCTTTTGAGT	GCAATTCCCT	AAAACCAGTA	CTCTAATAGT	TTTTCCCTAGA

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

326	TCACCTAGAT	CCTTTTAAAT	TAAAAATGAA	GTTTAAATC	AATCTAAAGT
	AGTGGATCTA	GCAAAATTTA	ATTTTACTT	CAAAATTTAG	TTAGATTTC
376	ATATATGAGT	AAACTTGGTC	TGACAGTTAC	CAATGCTTAA	TCAGTGAGGC
	TATATACTCA	TTTGAACCAG	ACTGTCAATG	GTTACGAATT	AGTCACTCCG
426	ACCTATCTCA	GGATCTGTC	TATTTCTGTC	ATCCATAGTT	GCCTGACTCC
	TGGATAGAGT	CGCTAGACAG	ATAAAGCAAG	TAGGTATCAA	CGGACTGAGG
476	CCGTCGTGTA	GATAACTACG	ATACGGGAGG	GCTTACCATC	TGGCCCCAGT
	GGCAGCACAT	CTATTGATGC	TATGCCCTCC	CGAATGGTAG	ACCGGGGTCA
526	GCTGCAATGA	TACCGCGAGA	CCCACGCTCA	CCGGCTCCAG	ATTATCAGC
	CGACGTTACT	ATGGCGCTCT	GGTGCGGAGT	GGCCGAGGTC	TAAATAGTCC
576	AATAAACCAG	CCAGCCGGAA	GGCCGAGCG	CAGAAGTGGT	CCTGCAACTT
	TTATTTGGTC	GGTCGGCCTT	CCC GGCTCGC	GTCTTCACCA	GGACGTTGAA
626	TATCCGCCCTC	CATCCAGTCT	ATTAAGTGT	GCCGGGAAGC	TAGAGTAAGT
	ATAGGCGGAG	GTAGGTCAGA	TAATTGACAA	CGGCCCTTCG	ATCTCATTTCA
676	AGTTCGCCAG	TTAATAGTTT	GCGCAACGTT	GTTGCCATTG	CTACAGGCAT
	TCAAGCGGTC	AATTATCAAA	CGCGTTGCAA	CAACGGTAAC	GATGTCCGTA

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

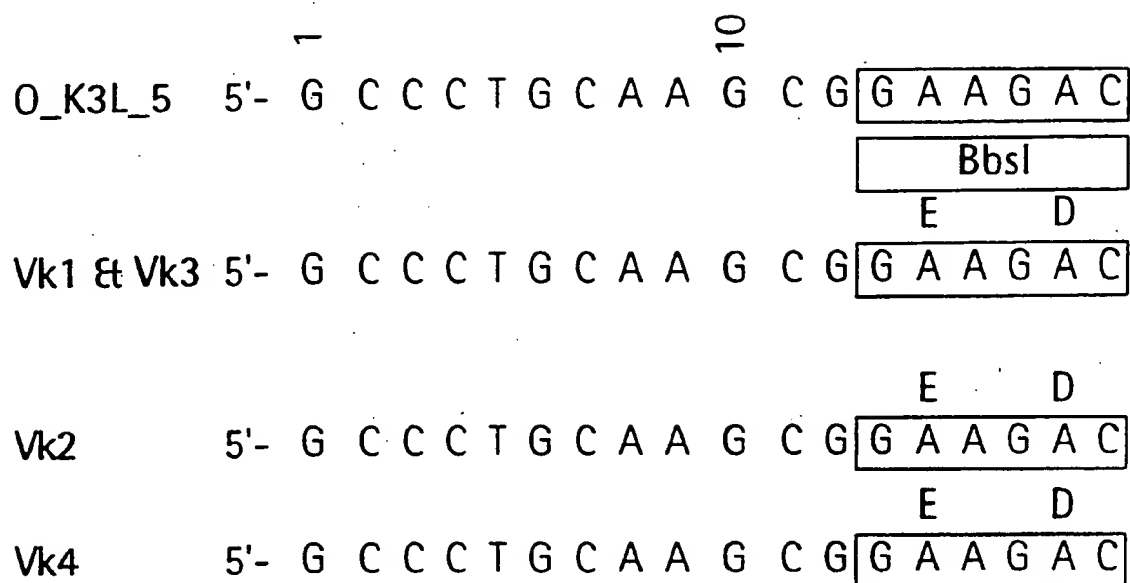
726	CGTGGTGTCA	CGCTCGTCGT	TTGGTATGGC	TTCATTACGC	TCCGGTTCCC
	GCACCACAGT	GCGAGCAGCA	AACCATACCG	AAGTAAGTCG	AGGCCAAGGG
776	AACGATCAAG	GCGAGTTACA	TGATCCCCCA	TGTTGTGCAA	AAAAGCGGTT
	TTGCTAGTTC	CGCTCAATGT	ACTAGGGGGT	ACAACACGTT	TTTTTCGCCAA
826	AGCTCCTTCG	GTCCCTCCGAT	CGTTGTCAGA	AGTAAGTTGG	CCGCAGTGTT
	TCGAGGAAGC	CAGGAGGCTA	GCAACAGTCT	TCATTCAACC	GGCGTCACAA
876	ATCACTCATG	GTTATGGCAG	CACTGCATAA	TTCTCTTACT	GTCATGCCAT
	TAGTGAGTAC	CAATACCGTC	GTGACGTATT	AAGAGAATGA	CAGTACGGTA
926	CCGTAAGATG	CTTTTCTGTG	ACTGGTGAGT	ACTCAACCAA	GTCATTCTGA
	GGCATTCTAC	GAAAGACAC	TGACCACTCA	TGAGTTGGTT	CAGTAAGACT
976	GAATAGTGT	TGCGGCGACC	GAGTTGCTCT	TGCCCGGCGT	CAATACGGGA
	CTTATCACAT	ACGCCGCTGG	CTCAACGAGA	ACGGGCCGCA	GTTATGCCCT
1026	TAATACCGCG	CCACATAGCA	GAAC TT TAAA	AGTGCTCATC	ATTGGAAAAC
	ATTATGGCGC	GGTGATCCGT	CTTGAAATTT	TCACGAGTAG	TAACCTTTTG
1076	GTTCTTCGGG	GCGAAAAC TC	TCAAGGATCT	TACCGCTGTT	GAGATCCAGT
	CAAGAAAGCCC	CGCTTTTGAG	AGTTCCTAGA	ATGGCGACAA	CTCTAGGTCA

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

1126	TCGATGTAAC	CCACTCGTGC	ACCCAACCTGA	TCTTCAGCAT	CTTTACTTT
	AGCTACATTG	GGTGAGCAG	TGGGTGACT	AGAAGTCGTA	GAAATGAAA
		BSSI	Eco57I		
		~~~~~	~~~~~		
1176	CACCAGCGTT	TCTGGGTGAG	CAAAACAGG	AAGCAAAAT	GCCGCAAAA
	GTGTCGCAA	AGACCCACTC	GTTTGTGTC	TTCCGTTTA	CGCGTTT
1226	AGGGAATAAG	GGGACACGG	AAATGTTGAA	TACTCATACT	CTTCTTTT
	TCCCTTATTC	CCGCTGTGCC	TTTACAACTT	ATGAGTATGA	GAAGGAAAA
1276	CAATATTATT	GAAGCATTTA	TCAGGGTTAT	TGTCTCATGA	GCGGATACAT
	GTTATAATAA	CTTCGTAAAT	AGTCCCAATA	ACAGAGTACT	CGCCTATGTA
		PstI	XhoI		
		~~~~~	~~~~~		
	EagI	BSSI	BbeI	AseI	BssHII
	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~
1326	ATTGAAATGT	ACTCGGCCGC	ACGAGCTGCA	GGGCCATTA	ATGGCTCGAG
	TAAACTTACA	TGAGCCGGCG	TGCTCGACGT	CCGCGGTAAT	TACCGAGCTC
	BssHII	BspEI	BsrGI		
	~~~~~	~~~~~	~~~~~		

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

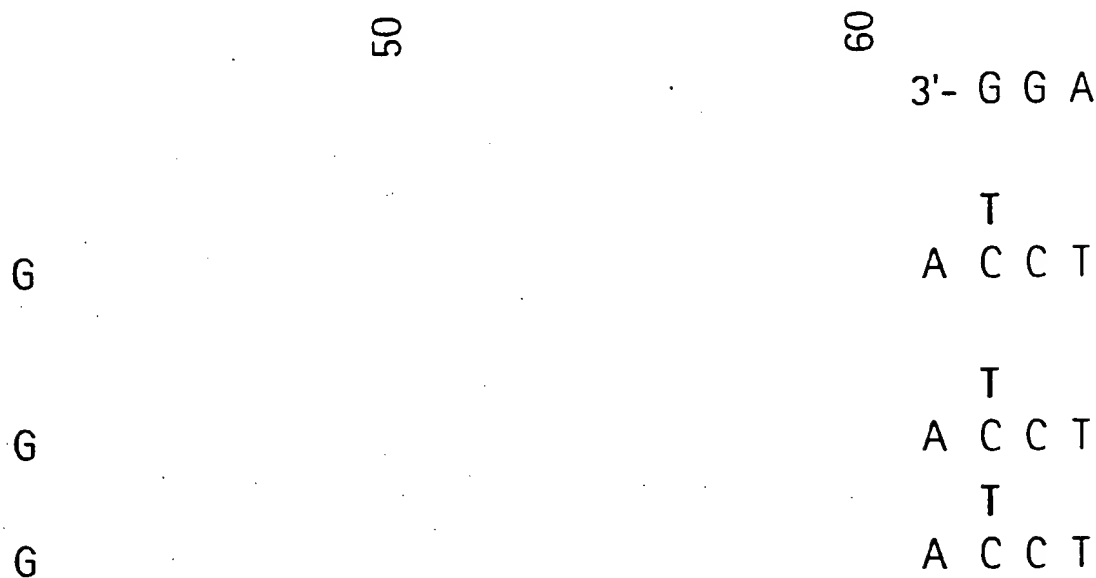
1376	CGCGCTTCAG	CGCTTTGTCT	TCCGGATGTA	CATGAAATT
	GCGCGAAGTC	GCGAAACAGA	AGGCCTACAT	GTACTTTTAA
	ECO57I	BbsI	~~~~~	~~~~~

Figure 37: Oligo and primer design for V<sub>κ</sub> CDR3 libraries

[illegible]

A	
C	
D	
E	
F	T T T
G	
H	C A T
I	
K	
L	C T T
M	A T G
N	
P	
Q	C A G
R	
S	
T	
V	
W	
Y	
80% Q	

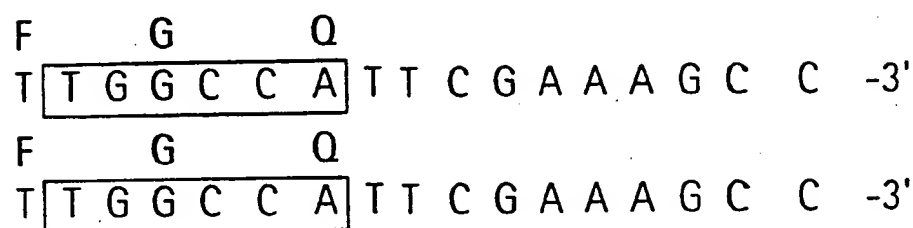


Figure 37: Oligo and primer design for V $\kappa$  CDR3 libraries

G C T			G C T		G C T
G A T	G A T	G A T	G A T		G A T
G A G			G A G		G A G
T T T			T T T		T T T
G G T	G G T	G G T	G G T		G G T
C A T			C A T		C A T
A T T			A T T		A T T
A A G			A A G		A A G
C T T			C T T		C T T
A T G			A T G		A T G
A A T	A A T	A A T	A A T		A A T
			C C T	C C T	C C T
C A G			C A G		C A G
C G T			C G T		C G T
T C T	T C T	T C T	T C T	T C T	T C T
A C T			A C T		A C T
G T T			G T T		G T T
T G G			T G G		T G G
T A T	T A T		T A T		T A T
50% Y			80% P		

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A A C C G G T A A G C T T T C G G -5' O\_K3L\_3



1                      10                      20

E       D       E       A       D

5'- CCTGCAAGCG GAAGAC GAAGC GGATT -

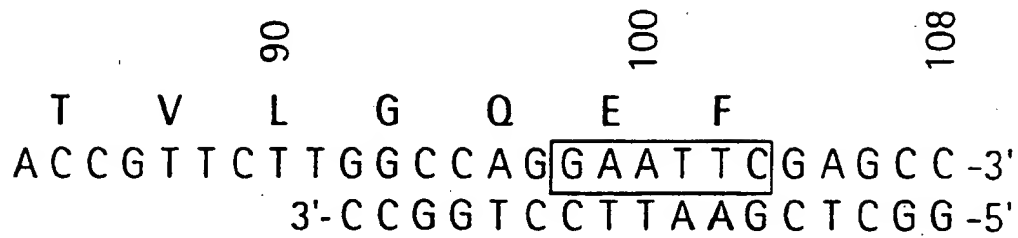
Figure 38: Oligo and primer design for V $\alpha$  CDR3 libraries

30					40		50			
Y	Y	C	Q	S			D			
-ATTATTGCCAGAGC							GAC			
					<div style="border: 1px solid black; padding: 2px;">A C D E F G H I K L M N P Q R S T V W Y</div>		<div style="border: 1px solid black; padding: 2px;">GCTGCT GATGAT GAGGAG TTTTTT GGTGGT CATCAT ATTATT AAGAAG CTTCTT ATGATG AATAAT CCTCCT CAGCAG CGTCGT TCTTCT ACTACT GTTGTT TATTAT</div>			

Figure 38: Oligo and primer design for VA CDR3 libraries

				60					70					80								
								G	G	G	T	K	L									
								G	G	C	G	G	C	A	C	G	A	A	G	T	T	A
-	G	C	T	gap	G	C	T	gap	G	C	T	G	C	T								
G	A	T	G	A	T	G	A	T	G	A	T	G	A	T								
G	A	G	G	A	G	G	A	G	G	A	G	G	A	G								
T	T	T	T	T	T	T	T	T	T	T	T	T	T	T								
G	G	T	G	G	T	G	G	T	G	G	T	G	G	T								
C	A	T	C	A	T	C	A	T	C	A	T	C	A	T								
A	T	T	A	T	T	A	T	T	A	T	T	A	T	T								
A	A	G	A	A	G	A	A	G	A	A	G	A	A	G								
C	T	T	C	T	T	C	T	T	C	T	T	C	T	T								
A	T	G	A	T	G	A	T	G	A	T	G	A	T	G								
A	A	T	A	A	T	A	A	T	A	A	T	A	A	T								
C	C	T	C	C	T	C	C	T	C	C	T	C	C	T								
C	A	G	C	A	G	C	A	G	C	A	G	C	A	G								
C	G	T	C	G	T	C	G	T	C	G	T	C	G	T								
T	C	T	T	C	T	T	C	T	T	C	T	T	C	T								
A	C	T	A	C	T	A	C	T	A	C	T	A	C	T								
G	T	T	G	T	T	G	T	T	G	T	T	G	T	T								
														T	G	G						
T	A	T	T	A	T	T	A	T	T	A	T	T	A	T								
18													19	Variability								
18	18												19	3.32E+05								
18	18	18											19	5.98E+06								
18	18	18	18										19	1.08E+08								

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Figure 38: Oligo and primer design for V $\lambda$  CDR3 libraries

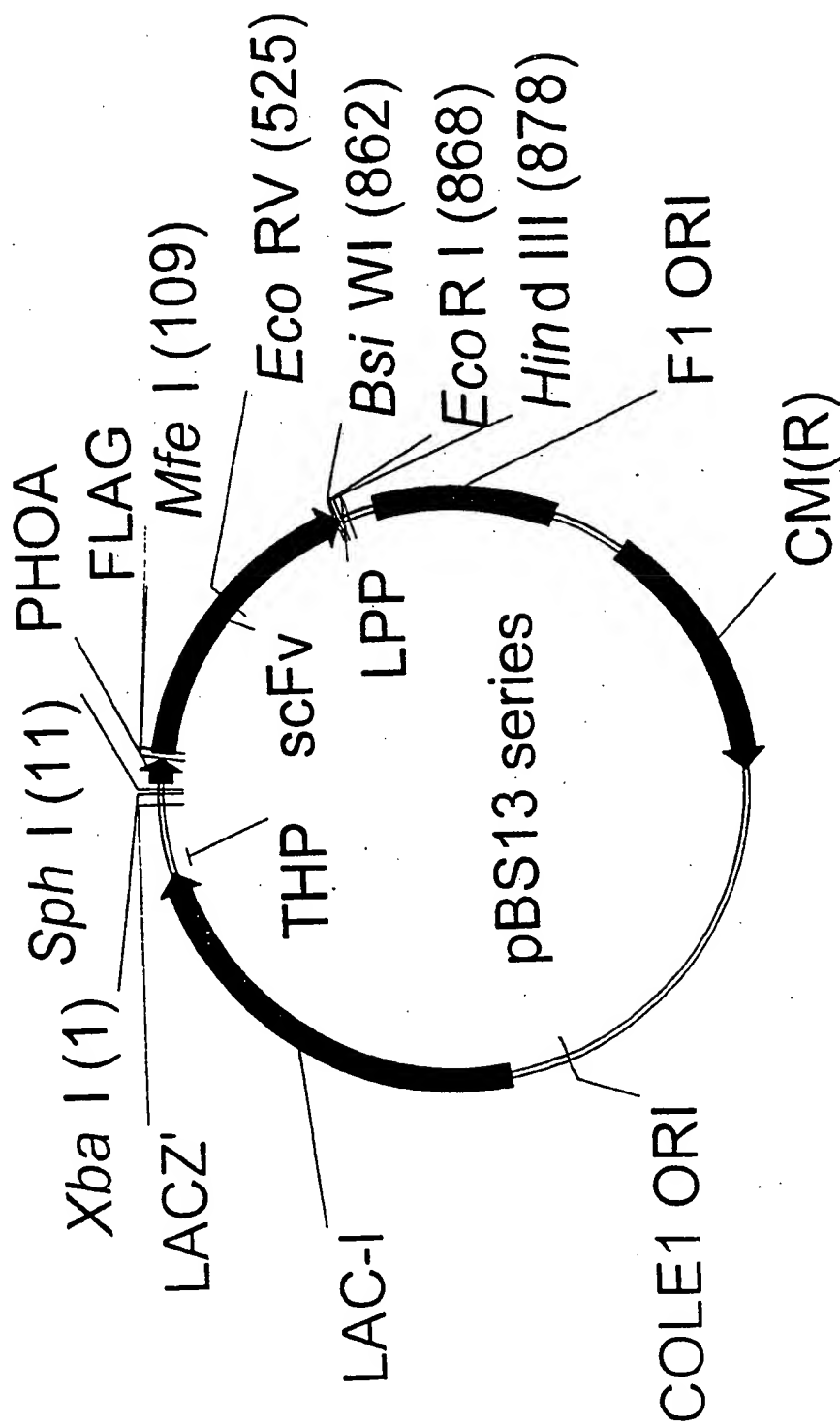


Figure 39: functional map of expression vector series pBS13

Figure 40: Expression data for HuCAL scFvs (pBS13, 30°C)

% soluble	$\kappa 1$	$\kappa 2$	$\kappa 3$	$\kappa 4$	$\lambda 1$	$\lambda 2$	$\lambda 3$
H1A	61%	58%	52%	42%	90%	61%	60%
H1B	39%	48%	66%	48%	47%	39%	36%
H2	47%	57%	46%	49%	37%	36%	45%
H3	85%	67%	76%	61%	80%	71%	83%
H4	69%	52%	51%	44%	45%	33%	42%
H5	49%	49%	46%	67%	54%	46%	47%
H6	90%	58%	54%	47%	45%	50%	51%

Total amount compared to H3 $\kappa 2$	$\kappa 1$	$\kappa 2$	$\kappa 3$	$\kappa 4$	$\lambda 1$	$\lambda 2$	$\lambda 3$
H1A	289%	94%	166%	272%	20%	150%	78%
H1B	219%	122%	89%	139%	117%	158%	101%
H2	186%	223%	208%	182%	126%	60%	97%
H3	50%		71%	54%	59%	130%	47%
H4	37%	55%	60%	77%	195%	107%	251%
H5	98%	201%	167%	83%	93%	128%	115%
H6	65%	117%	89%	109%	299%	215%	278%



Figure 40: Expression data for HuCAL scFvs (p8S13, 30°C)

Soluble amount compared to H3κ2	κ1	κ2	κ3	κ4	λ1	λ2	λ3
H1A	191%	88%	121%	122%	26%	211%	76%
H1B	124%	95%	83%	107%	79%	142%	59%
H2	126%	204%	139%	130%	66%	50%	70%
H3	63%	-	81%	49%	69%	143%	61%
H4	40%	47%	49%	54%	95%	55%	125%
H5	69%	158%	116%	80%	72%	84%	84%
H6	85%	122%	87%	77%	162%	162%	212%
	McPC						
soluble	38%						
%H3κ2 total	117%						
%H3κ2 soluble	69%						